## STIC Search Report Biotech-Chem Library



## STIC Database Tracking Number 119952

From: Paul Schulwitz

Location: Biotech-Chem Library

REM-1A65

Phone: (571)272-2527

paul.schulwitz@uspto.gov

Search Notes

**2681** :JinU hA

Examiner Schultz,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Case Serial Number: 10/001844

Wednesday, April 21, 2004

Location: REM-2D18/2C18

TO: James Schultz

Paul Schulwitz Technical Information Specialist STIC Biotech/Chem Library (S11)272-2527



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Human Acetylcholin Human ALDHS allele Human/mouse C/EBP Human Sonic hedgeh Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bloavalibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss. Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:43. ALIGNMENTS AAS96144 ABA99313 ABA02229 AAF27039 멾 RESULT 1 AAF27039/c ID AAF27039 standard, DNA; 38 30-MAR-2001 (first entry) 30 12 AAF27039; 

26-MAY-2000; 2000WO-US014741. 01-JUN-1999; 99US-0137011P. 13-AUG-1999; 99US-0149016P. (BIOJ ) BIOGEN INC.

WO200073337-A1.

07-DEC-2000.

Homo sapiens.

Synthetic.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues. Garber Pepinsky RB, Taylor F, WPI; 2001-049927/06.

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant sonic hedgehog protein by indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of a defining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the correct or the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders.

Command the hedgehog polymer conjugates will cancers. In particular, they may be used to prevent preventing or ameliorate neurological disorders (e.g., Parkinson's disease, Allentimet's disease), age-associated neurological disease, neurological injury and trauma; immunological diseases of the nervous system (e.g., multiple collerosis); stroke; and malignant gliomas, medulloblastomas and neuroectodermal tumours. The modifications made to the hedgehog protein may result in increased half-life, altered tissue distribution (such as

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an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from protectly degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human solution primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention
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Sequence 38 BP; 8 A; 11 C; 9 G; 10 T; 0 U; 0 Other;

Gaps ö Ouery Match

8.5%; Score 36.4; DB 1; Length 38;
Best Local Similarity 97.4%; Pred. No. 0.031;
Matches 37; Conservative 0; Mismatches 1; Indels 162 GACTGGGTGTACTACGAGTCCAAGGCACATATCCACTG 199 ð

38 GACTGGGTGTACTACGAGTGCAAGGCACATATCCACTG 1

AAF27025 standard; DNA; 49 BP. AAF27025; 

30-MAR-2001 (first entry)

Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:29.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; siroke; multiple sclerosis; multigant glioma; medulloblastoma; neuroectodermal tumour;

Homo sapiens Synthetic. WO200073337-A1.

07-DEC-2000.

26-MAY-2000; 2000WO-US014741.

01-JUN-1999; 99US-0137011P.

(BIOJ ) BIOGEN INC.

Pepinsky RB, Taylor F,

Garber E;

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.

Example 2; Page 67; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bicavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lyshne residues of the hedgehog protein. The hedgehog protein used in the conjugate may be a wild-type or mutant sonic hedgehog (Shh), Indian hedgehog (Lhh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of defining and mapping functionally important regions of a protein by

modifying accessible amino acid side chains, and determining the effect
the position and/or type of modification have on the activity of the
protein. The hedgehog polymer conjugates may be used in the management of
c various medical conditions including various neurological disorders,
inflammatory and autoimmune diseases, and cancers. In particular, they
any be used to prevent preventing or ameliorate neurodegenerative
cd disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's
disease); age-associated neurological disease; neurological injury and
traums; immunological diseases of the nervous system (e.g., multiple
cc traums; immunological diseases of the nervous system (e.g., multiple
cc reaums; immunological diseases of the nervous system (e.g., multiple
cc neuroectodermal tumours. The modifications made to the hedgehog protein
cc neuroectodermal tumours in the modifications made to the hedgehog protein
cc nay result in increased half-life, alterature for longer periods of
c time), increased stability to stay in the vasculature for longer periods of
ctime), increased stability in solution, protection from proteolytic
degradation, or reduced immunogenicity. In particular, the ability to
ce remain in the vasculature for prolonged periods may allow a hedgehog
cc protein of the invention to cross the blood-brain barrier, and an
cincreased thermal stability would be an advantage when formulating the
chedgehog protein in powder form. The present sequence represents a human
cc increased dedepend mutagenic primer used in an exemplification of the

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Sequence 49 BP; 8 A; 18 C; 9 G; 14 T; 0 U; 0 Other;

.. 0 8.5%; Score 36; DB 1; Length 49; 88.6%; Pred. No. 0.067; tive 0; Mismatches 5; Indels Query Match Best Local Similarity 88.6 Matches 39, Conservative

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AAF27038 standard; DNA; 39 BP. AAF27038/c RESULT 3

AAF27038;

30-MAR-2001 (first entry)

Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:42.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bloavallibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss.

Homo sapiens. Synthetic.

WO200073337-A1. 07-DEC-2000. 26-MAY-2000; 2000WO-US014741.

99US-0137011P. 99US-0149016P. 01-JUN-1999; 13-AUG-1999; 

Pepinsky RB, Taylor F,

BIOJ ) BIOGEN INC.

Garber E;

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyalkylene conjugated to a non-naturally-occurring polymer comprising a polyalkylene glycol group, with the proviso that the polymer is not conjugated to the N-terminus, or to lysine residues of the hedgehog protein. The hedgehog protein. The hedgehog protein in the conjugate may be a wild-type or mutant sonic hedgehog (Shh), indian hedgehog (Ihn) or besert hedgehog protein. The mapping functionally important regions of a protein by modifiving accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, inflammatory and autoinmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders (e.g., Parkinson's disease, Huntington's disease, Alzheimer's disease, associated neurological diseases, neurological injury and traumay immunological diseases, medulioblastomas and cancercine in increased half-life, altered tissue distribution (such as an improved ability to stay in the vasculature for longer periods of time), increased stability in solution, protection from protectory can improved ability to stay in the vasculature for prolonged periods may allow a hedgehog conjugated in powder form in particular, the ability to deay in the vasculature for prolonged periods may allow a hedgehog protein in powder form. The present sequence represents a human conjugation of the hedgehog mutagenic primer used in an exemplification of the

Sequence 39 BP; 7 A; 12 C; 13 G; 7 T; 0 U; 0 Other;

Gaps ; Score 35.8; DB 1; Length 39; Pred. No. 0.043; 2; Indels 97 CCACGTCTGACCGCGACCGCAGCAAGTACGGCATGCTGG 135 н 39 CCACGTCTGACCGCGATCGCTGCAAGTACGGCATGCTGG 0; Mismatches 8.4%; Local Similarity 94.9 les 37, Conservative Query Match Matches ઠ 셤

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AAF27037/c ID AAF27037 standard; DNA; 37 BP (first entry) 30-MAR-2001 AAF27037; RESULT 

Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:41.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bicavailbbility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; parkinson's disease; Huntington's disease; halzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss

Homo sapiens. Synthetic.

07-DEC-2000,

WO200073337-A1.

26-MAY-2000; 2000WO-US014741.

99US-0137011P. 1999;

99US-0149016P 13-AUG-1999;

(BIOJ ) BIOGEN INC.

M Garber 7 Taylor Pepinsky RB,

WPI; 2001-049927/06.

useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine protein, Modified hedgehog

Example 6; Page 77; 157pp; English

The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyarlylene corpiugated to a non-naturally-occurring polymer comprising a polyarlylene cyptoel group, with the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant Sonic hedgehog corputation used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or beart hedgehog protein. The hedgehog (Ihh) or beart hedgehog protein, or may be cefining and mapping functionally important regions of a protein by condition and/or type of modification have on the activity of the postion and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the amanagement of various medical conditions including various neurological diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disease); inflammatory and autoimmune diseases, Huntington's disease, Alzheimer's disease, inflammatory and autoimmune diseases, mediloblastomas and trauma; immunological diseases, Huntington's disease, Alzheimer's disease, associated neurological diseases, incurlogical injury and traumy immunological diseases of the nervous system (e.g., multiple calerosis); stroke, and malignant gliomas, mediloblastomas and neurococodermal tumours. The modifications made to the hedgehog protein captored immunogenicity. In particular, the ability to stay in the vasculature for longer periods may remain in the vasculature for prolonged periods may allow a hedgehog cremain in the vasculature form protection from the vasculature form protection from the invantion to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating them to may a non-independent of the invantion to cross the blood-brain and advantage when formulating in an exemplificat

Sequence 37 BP; 6 A; 10 C; 12 G; 9 T; 0 U; 0 Other;

Gaps ö 7.9%; Score 33.8; DB 1; Length 37; 4.6%; Pred. No. 0.099; ve 0; Mismatches 2; Indels 94.68; 35; Conservative Query Match Best Local Similarity

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Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:45. 041/c AAF27041 standard; DNA; 35 (first entry) 30-MAR-2001 AAF27041; AAF27041/ RESULT 5 XXXEXEXEXEX

bioavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease; Alzheimer's disease; neurological injury; stroke; multiple sclerosis; malignant glioma; medulloblastoma; neuroectodermal tumour; Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group;

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26-MAY-2000; 2000WO-US014741
                                       Pepinsky RB, Taylor F,
mutagenic primer; ss
                                 (BIOJ ) BIOGEN INC.
            WO200073337-A1
     Homo sapiens.
                           01-JUN-1999;
                 07-DEC-2000.
                                                                                                                                          invention
       Synthetic
                                                         residues
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The invention relates to novel polymer conjugates of hedgehog proteins which have increased bioavailability. The hedgehog proteins are which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polymer of glycol group, with the proviso that the polymer is not conjugated to the conjugated to a non-naturally-occurring polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant Sonic hedgehog or protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Ihh) or Desert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of a hedgehog fusion and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological disorders, infammatory and autoimmune diseases, and cancers. In particular, they may be used to prevent preventing or ameliorate neurodegenerative disorders (e.g., Parkinson's disease, huntington's disease, Alzheimer's disease); age-associated neurological disease, huntington's disease, Alzheimer's calescals); stroke; and malignant gliones, meduloblastoms and consumary immunological diseases of the nervous system (e.g., multiple or stay in the vasculature for longer periods of time), increased stability in solution, protection from proteclytic degradation, or reduced immunogenicity. In particular, the ability to remain in the vasculature for prolonged periods may allow a hedgehog protein of the invention to crose the blood-brain harrier and an
                                                                                           Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein of the invention to cross the blood-brain barrier, and an increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human sonic hedgehog mutagenic primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; Page 77; 157pp; English.
WPI; 2001-049927/06
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ö Score 33.4; DB 1; Length 35; Pred. No. 0.1; 0; Mismatches 1; Indels Sequence 35 BP; 8 A; 15 C; 9 G; 3 T; 0 U; 0 Other; 7.8%; 97.1%; 34; Conservative Similarity Query Match Best Local S Matches 34

139 GCCTGGCGGTGGAGGCCGGCTTCGACTGGGTGTAC 173 35 GCCTGGCGGTGGAGGCCTGCTTCGACTGGGTGTAC

d 8

AAF27040 standard; DNA; 37 BP AAF27040/c RESULT 6

AAF27040; 

(first entry) 30-MAR-2001 Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:44.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bloavailibility; formulation; neurological disorder; inflammatory disorder; autoimmune disorder; cancer; neurodegenerative disorder; parkinson's disease; Huntington's disease, Alzheimer's disease; neurological injury; stroke; multiple sclerosis; multignant glioma; medulloblastoma; neuroectodermal tumour; mutagenic primer; ss.

Homo sapiens Synthetic. WO200073337-A1

Garber E;

99US-0137011P

07-DEC-2000.

26-MAY-2000; 2000WO-US014741.

99US-0137011P. 99US-0149016P 01-JUN-1999; 13-AUG-1999;

(BIOJ ) BIOGEN INC.

ü Garber Pepinsky RB, Taylor F,

WPI; 2001-049927/06.

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.

Example 6; Page 77; 157pp; English.

The invention relates to novel polymer conjugates of hedgehog proteins are which have increased bioavailability. The hedgehog proteins are which have increased bioavailability. The hedgehog proteins are conjugated to a non-naturally-occurring polymer comprising a polyarlylene of glycol group, with the proviso that the polymer is not conjugated to the proviso that the polymer is not conjugated to the proviso that the polymer is not conjugated to the protein used in the conjugate may be a wild-type or mutant Sonic hedgehog (Shh), Indian hedgehog (Thh) or besert hedgehog (Dhh) protein. The hedgehog (Ihh) or besert hedgehog (Dhh) protein, or may be a hedgehog fusion protein. The invention also relates to methods of cefining and mapping functionally important regions of a protein by modifying accessible amino acid side chains, and determining the effect the position and/or type of modification have on the activity of the protein. The hedgehog polymer conjugates may be used in the management of various medical conditions including various neurological diseases, and cancers. In particular, they warious he used to prevent preventing or ameliocate neurodegenerative disorders (e.g., Parkinson's disease, neurological disease, Alzheimer's disease, neurological disease, neurological disease, insurological injury and creama; immunological diseases of the nervous system (e.g., multiple colerosis); stroke; and malignant gluomas, medulloblastomas and conscretodermal tumours. The modifications made to the hedgehog protein any presult in increased thalf-life, altered tissue distribution (such as an improved ability to stay in the vasculature for prolonged periods may allow a hedgehog conceased tebrility in solution, protection from the vasculature for prolonged periods may allow a hedgehog contens in powder form. The present sequence represents a human considerion made of the numberion of the num nvention

Sequence 37 BP; 7 A; 8 C; 13 G; 9 T; 0 U; 0 Other;

Query Match

7.6%; Score 32.2; DB 1; Length 37;

ABT03768/c

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The sequences given in AAQ91654-57 are primers which were used to amplify a sequence which encodes a human sonic hedgehog protein, homologous to a brosophila hedgehog protein (AAR77337). The human sequence was isolated by screening of human genome DNA by nested polymerase chain reaction using these primers, followed by use of a clone to screen a human fetal lung 5'-stretch plus cDNA library in phage lambda-gt10. A clone has been primers SHHF (AAQ91654) and SHH (AAQ91655), to give clone SHHP1. A 2.5-kb ECORI CA repeat fragment is amplified using primers SHHCAR (AAQ91656) and SHHPAR (AAQ91657). Probes and primers derived from the sonic hadgehog sequence may be used as diagnostic agents for neuromuscular, autonomic or central nervous system disorders, and the gene may also be used in gene therapy. Antibodies generated from the encoded protein may be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sonic hedgehog; SHH gene; HH gene; tumorigeneais; oncogeneais; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hedgehog-like protein(s) and nucleic acid(s) encoding them - useful treat degenerative nervous system disorder(s) and in gene therapy.
            Human, sonic hedgehog gene; nested polymerase chain reaction; PCR; fetal lung; probe; primer; diagnostic; nervous system disorder; gene therapy; antibody; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mutated sonic hedgehog (SHH) gene exon 2 PCR primer.
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100.0%; Pred. No. 3.8;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                    (HARD ) HARVARD COLLEGE.
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY.
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94US-00356060
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                                                                                                                                        WO9518856-A1
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14-DEC-1994;
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                                                                                                 Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for determining the presence of neoplastic molecular markers in a host, involving the use of neoplastic molecular marker specific reagents to detect such markers and analysing the array of reagents, allowing the identification of the neoplastic disease present. This can be used to determine the best treatment for cancers, in particular neural cell, lung and prostate tumours. The present sequence is a PCR primer useful for detecting the coding sequences of markers of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Determining the presence of neoplastic molecular markers, by identifying the presence of markers in host test sample using array of neoplastic molecular marker specific reagents and analyzing the array of the
                                                                                                                                                                                                                                                                                                                                                         Human, cancer; neoplastic disease; tumour specific marker; cytostatic; transcription factor; PCR; primer; ss.
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100.0%; Pred. No. 1.2;
iive 0; Mismatches 0; Indels
                       Indels
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                       <u>ښ</u>
                                                         38 CGAAGATGGCCACCACTCAGAGGAGTCTCTGCACTAC 74
                                                                                       CGAAGATGGCCACCACTCATGCGAGTCTCTGCACTAC 1
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  Pred. No. 0.21;
0; Mismatches
                                                                                                                                                                                                                                                                                                                         Human SHH gene PCR primer SEQ ID NO: 289.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 19; 41pp; English.
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91.9%;
                                                                                                                                                                                                   ABT03768 standard; DNA; 27
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  Best Local Similarity 91.9
Matches 34; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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4AQ91654 RESULT

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This human sonic hedgehog (SHH) gene exon 2-specific primer was used with another exon 2-specific primer (see AAV18406) in a PCR using DNA from Muman bacterial artificial chromosome (BAC) DNA pools. Only pools comprising a BAC that contains the sequence tag defined by the primer pair will yield an amplification product. The process was continued until as ingle positive BAC was identified. The positive clone, BAC270A17, was dispessed with restriction enzymes and ligated into vectorette linkers. Mutations (see AAV18403 and AAV18404) have been identified in the SHH gene in Muman cancers. The mutated SHH genes and the encoded polypeptides (see AAW48735 and AAW48736) can be used in methods for the treatment and diagnosis of cancer and other diseases involving cell proliferation or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, Sonic hedgehog, Shh, morphogenic signal; neuron; embryonic patterning; cell culture; cell differentiation; ischaemia; cell proliferative disorder; intracerebral grafting; Huntington's chorea; neurological disorder; Alahaimer's disease; Parkinson's disease; amyotrophic lateral sclerosis; ALS; multiple sclerosis; PCR primer; ss.
                                                                                                                                                                                                          New nucleic acid encoding oncogenic human hedgehog protein - useful for, e.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Sonic hedgehog (Shh) gene amplifying forward PCR primer SHHF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.6%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 3.8;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                            Example; Page 23; 47pp; English.
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94US-00356060.
95US-00435093.
95US-00460900.
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                                               97WO-US020227
                                                                            96US-00748591
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                                                                                                             (REGC ) UNIV CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             differentiation
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05-JUN-1995;
05-JUN-1995;
                                               12-NOV-1997;
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                 22-MAY-1998.
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                                                                                                                                               Epstein E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD10171;
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Gaps

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The present invention relates to assay for screening compounds that corporate or inhibit binding of hedgehog polypeptide to naturally concurring patched receptor. The hedgehog proteins comprise morphogenic signals produced by embryonic patcarning centres, and are involved in the formation and maintenance of ordered spatial arrangements of differentiated tissues in vertebrates, both adult and embryonic. The proteins can be used to generate and/or maintain a marray of different corporate tissues both in vitro and in vivo. The invention also relates to ammanian cell (e.g. neuron, testicular cell) responsive to hedgehog induction. Hedgehog agonists and anicagomists can be used in cell culture techniques to enhance survival and maintenance of neurons and various vertebrate organogenic patchways. The hedgehog gene is useful in determining whether a patient is at the risk of disorder characterised by unwanted cell proliferation or aberrant control of differentiation. The ceptangon proteins or mimetics can be used to induce foctal neurons especially neuronal stem cells in intracerebral grafting. The protein or its mimetic can be used in the treatment of neurological conditions e.g. cinjury to nervous system, isochaemia resulting from Erroke, Alzheimer's condisease, Parkinson's disease, Hunthington's chorea, amyotrophic lateral conditions c
                                                                                                                                                     Screening compounds that potentiate or inhibit binding of hedgehog polypeptide to naturally occurring patched receptor, comprises contacting polypeptide with receptor and test compound, and detecting change in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.6%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 3.8;
ive 0; Mismatches 0; Indels
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                                                                    Mcmahon AP
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD. (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Shh DNA amplifying primer SHHF5'.
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                                                                                                                                                                                                                                                                            Example 5; Col 98; 127pp; English
                                                                  Ingham PW,
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94US-00356060.
95US-00435093.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH76132 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100.
                                                                    Tabin CJ,
                                                                                                                  WPI; 2001-440859/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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14-DEC-1994;
04-MAY-1995;
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                                                                    Marigo V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH76132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                binding
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Matches
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AAH76132
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for treating diseases

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PCR primers AAC87097-98 were used to amplify cDNA encoding a hedgehog related-protein. The specification describes a sonic hedgehog protein (ShN), a desernt hedgehog protein (DhN), and an indian hedgehog protein (IN). The hedgehog polynuclectides are useful in diagnostic, in antisense therapy and in therapeutic assays for detecting and treating disorders involving, e.g., aberrant expression of vertebrate hedgehog completes involving, e.g., aberrant expression of vertebrate hedgehog completes involving, e.g., aberrant expression of vertebrate hedgehog conditions deriving from acute, subacute, or chronic injury to the conditions deriving from acute, subacute, or chronic injury, vasal injury conditions system, including traumatic injury, chemical injury, vasal injury conditions from stroke), together with infectious/inflammatory and induced injury, aging of the nervous system including Alzheimer's disease, chronic neurodegenerative diseases of the nervous system, including Parkinson's disease, Huntington's chorea, amylotrophic lateral sclerosis, spinocerebellar degenerations, and chronic immunological diseases of the nervous system or affecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sonic hedgehog; Shh; desert hedgehog; Dhh; Indian hedgehog; Ihh; antiparkinsonian; antiarrhythmic; neuroprotective; anticonvulsant; cytostatic; noctropic; spermatogenesis; peripheral nervous system; central nervous system; Alzheimer's disease; Parkinson's disease; Huntington's disease; arrhythmia; nerve degeneration; multiple sclerosis; immunological disorder; neoplastic; hyperplastic; PCR primer; ss.
               Polynucleotides encoding hedgehog proteins, useful for treating diseases of nervous system such as Alzheimer's disease, Parkinson's disease, Huntington's chorea, amylotrophic lateral sclerosis, multiple sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.6%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 3.8;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human sonic hedgehog (Shh) PCR primer SHHF SEQ ID NO:43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HARD ) HARVARD COLLEGE.
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24 ACCGAGGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACCGAGGCTGGGACGAAGATGGC 24
                                                                                         Example 5; Col 86; 119pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93US-00176427.
94US-00356060.
95US-00435093.
95US-00462386.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.
nes 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ingham PW, Mcmahon AP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6384192-B1.
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14-DEC-1994;
04-MAY-1995;
05-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                   The invention relates to nucleic acids encoding hedgehog proteins selected from sonic hedgehog (Shh), indian hedgehog (Ihh), desert hedgehog (Dh) polypeptides. The hedgehog genes are involved in the formation of ordered spatial arrangements of differentiated tissue in vertebrates. The nucleic acid sequences are useful for producing hedgehog proteins, used for promoting differentiation of, or survival of differentiation on unronal cells, and for promoting proliferation, survival or differentiation of mesenchymal, endodermal or ectodermal tissue, particularly chondrocytes, or testicular germ line cells. Sequences AAH76132-133 represent PCR primers for amplifying a human Shh DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hedgehog related-protein, sonic hedgehog protein, Shh; ischemia; stroke; desert hedgehog protein, Dhh; indian hedgehog protein, Ihh; neuron; neurological condition; nervous system injury; tumour-induced injury; aging; Alzheimer's disease; chronic neurodegenerative disease; barkinson's disease; thronic neurodegenerative disease; spinocease; thuntington's chorea; amylotrophic lateral sclerosis; spinoceachelar degeneration; chronic immunological disease; multiple sclerosis; PCR primer; ss.
                                                                                                                                                          Novel nucleic acid encoding a hedgehog polypeptide, used to produce the polypeptide, which is used to promote proliferation, survival, and/or differentiation of neuronal and mesodermal tissue.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer for cDNA encoding human sonic hedgehog protein (Shh).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
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(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                  IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24 ACCGAGGGTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ACCGAGGCTGGGACGAAGATGGC 24
                                                                                       Tabin CJ;
                                                                                                                                                                                                                                    Example 5; Col 88; 118pp; English.
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94US-00356060.
95US-00435093.
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95US-00462386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAC87097 standard; DNA; 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.
                                                                                         Ingham PW, Mcmahon AP,
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                                  (HARD ) HARVARD COLLEGE
                                                                                                                        WPI; 2001-456723/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-079847/09
05-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-DEC-1994;
04-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 12
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New vertebrate hedgehog-related proteins, useful e.g. for promoting differentiation, survival and proliferation of cells, e.g. for treating neurodegeneration. 

The present invention describes an isolated and/or recombinant configuration of comprising a hedgehog (hh) amino acid (aa) sequence polypeptide (I) comprising a hedgehog (hh) amino acid (aa) sequence concoded by a nucleic acid (II) that hybridizes under stringent conditions to 1 of 6 sequences (see ABN87544, and ABN87546 to ABN87550). (I) binds to a natural patched receptor. Specifically claimed example of (I) are concrepted neuroprotective, anticonvulsant, antiarrhythmic and cytostatic activities. (I) induces the expression of the BMP-2 and -4 genes, and of the Howa gene. (I) and be used: (i) to promete differentiation of courconal cells and survival of the differentiation of courconal cells and survival of the differentiation of copaminergic or motor neurons, proliferation of chondrocytes, and companiergic or motor neurons, proliferation of chondrocytes, and concordermal cells, either in cell cultures (particularly for preparation of transplants) or therapeutically; (ii) for detecting loss of response, in tissues or, to hh proteins, (iii) in drug screening (to identify (art) agonists, useful e.g. for inhibition of spermatogenesis); and (iv) correct collation of cognate receptors. (I) may be used therapeutically to treat e.g. injuries/defects in the central or peripheral nervous systems, e.g. multiple sclerosis, neoplastic and hyperplastic concorder system, e.g. multiple sclerosis, neoplastic and hyperplastic concorder system, e.g. multiple sclerosis, neoplastic or promote attachment of prostheses. The present sequence represents a PCR primer for human sonic concorder invention in the exemplification of the present Example 5; Col 88; 116pp; English.

Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

Query Match 5.6%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 3.8; Matches 24; Conservative 0; Mismatches 0; Indels 24 ACCGAGGCTGGGACGAAGATGGC 47 1 ACCGAGGCTGGGACGAAGATGGC 24 Бb

0; Gaps

ADA26284 standard; DNA; 24 BP. RESULT 14 ADA26284

20-NOV-2003 (first entry)

ADA26284;

Human Sonic hedgehog (Shh) cDNA PCR primer #1.

Human; PCR; ss; Sonic hedgehog; Shh; neuronal cell; skeletogenesis; chrondrogenesis; osteogenesis; degenerative disorder; nervous system; neuronal cell death; neural cell; neuromuscular disorder; acrosta autonomic disorder; central nervous system disorder; anoxia; ischaemia; peripheral nervous system disorder; acrobycardia; atrial cardiac arrhythmia; striated heart; stem cell development; digestive tract; liver; multiple sclerosis; primer.

Homo sapiens

20-MAR-2003.

JS2003054437-A1.

20-OCT-1997;

97US-00954771.

93US-00176427. 94US-00356060. 95US-00435093. 30-DEC-1993; 14-DEC-1994; 04-MAY-1995; 

95US-00462386 05-JUN-1995;

(INGH/) INGHAM P W. (MCMA/) MCMAHON A P. (TABI/) TABIN C J.

Tabin CJ; Ingham PW, Mcmahon AP,

WPI; 2003-555377/52

Modulating growth, differentiation or survival of a cell, useful for treating a degenerative disorder of the nervous system characterized by neuronal cell death, comprises contacting the cell with a hedgehog polypeptide.

Example 5; Page 48; 121pp; English.

The invention relates to a method for modulating growth, differentiation or survival of a cell, comprising contacting the cell with a hedgehog polypeptide. The invention also relates to methods for inducing a cell to differentiate to a neuronal cell phenotype comprising contacting the cell carget tissue of a hedgehog polypeptide to cause chrondrogenesis and/or osteogenesis in the target tissue and treating a degenerative disorder of the nervous system characterised by neuronal cell death, comprising cadministerings a hedgehog polypeptide causing prolonged survival of neural cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptides are useful for treating a degenerative disorder of cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptides are useful for treating adegenerative disorder of the nervous system characterised by neuronal cell death, including concerns accordance of hedgenoration accordance or central nervous system disorders, precipied and secondary autonomic or central nervous system disorders, precipied and secondary autonomic or central nervous system disorders including disorders affecting from anoxia, ischaemia or trauma and neuronal degeneration associated with a natural aging process. The call sorders affecting peripheral nervoe innervation of smooth muscle and peripheral nerve damage, for treating peripheral nerve damage, for treating neoplastic or hyperplastic and peripheral nerve damage, for treating neoplastic or hyperplastic crasping for the development of stem cells of the heart, in nerve prostheses for repairing central carponating the heart, in nerve prostheses for repairing central corpusible for the formation of the digestive tract, liver and other corpusion of the digestive race, liver and other corpusions. This sequence represents a PCR primer used to amplify condition the human Sonic hedgehog (Shh) polypeptide. 

Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

. 0 Match 5.6%; Score 24; DB 1; Length 24; Local Similarity 100.0%; Pred. No. 3.8; see 24; Conservative 0; Mismatches 0; Indels Query Match Best Loca Matches

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Gaps

g

RESULT 15 ADD25290

ADD25290 standard; DNA; 24 BP

ADD25290;

15-JAN-2004 (first entry)

Human Sonic hedgehog PCR primer #1.

hedgehog; patched receptor; spermatogenesis inhibition; ovary function inhibition; embryogenesis; differential tissue maintenance; ss; PCR; primer; human.

Homo sapiens.

US6576237-B1

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The invention relates to an isolated antibody (I) which is immunoreactive with a hedgehog polypeptide (II) that binds to a patched receptor, where (II) is encoded by uncleic acid which hybridise to a fully defined vertebrate hedgehog (hh) protein. (I) is useful as a hedgehog antagonist by blocking action of naturally occurring hedgehog protein, and therefore for inhibiting spermatogenesis. (I) is also useful for inhibiting normal ovarian function. (I) is useful for inhibiting normal ovarian function. (I) is useful for blocking the action of one or more hedgehog proteins and allows the study of the role of these proteins e.g., embryogenesis and/or maintenance of differential tissue succeed to evaluate the abundance and pattern of expression of the hedgehog protein abundance and pattern of expression of the hedgehog protein to detect and evaluate hedgehog procedure. The present sequence represents as a part of clinical testing procedure. The present sequence represents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, cell differentiation, Desert hedgehog, Dhh, Sonic hedgehog; shh, Indian hedgehog; Ihh; skeletogenesis; degenerative disorder; ischaemia; Alzheimer's disease; Parkinson's disease; parkinson's disease; amortophic lateral sclerosis; Huntington's disease; multiple sclerosis; Pick's disease; aging process; trauma; anoxia; antisense gene therapy; neuroprotective; anticonvulsant;
                                                                                                                                                                                                                                                                                                                                 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                         Novel isolated antibody which is immunoreactive with a vertebrate hedgehog protein sequence that binds with patched receptor, useful blocking action of naturally occurring hedgehog protein, and for
                                                                                                                                                                                                                                       Marti-Gorostiza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24; DB 1; Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                       Bumcrot DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.8;
                                                                                                                                                                                 (HARD ) HARVARD COLLEGE.
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 100.0%; Pred. No. 3.8 nes 24; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; SEQ ID NO 43; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24 ACCGAGGCTGGGACGAAGATGGC 47
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                                                                                       93US-00176427.
94US-00356060.
95US-00435093.
95US-00460900.
                                                      16-AUG-2000; 2000US-00639695.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.6%;
                                                                                                                                                                                                                                                                                                                                                                    inhibiting spermatogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAD62117 standard; DNA; 24
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                                                                                                                                                                                                                                       Ingham PW, Mcmahon AP,
                                                                                                                                                                                                                                                                          WPI; 2003-799823/75.
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                                                                                       30-DEC-1993;
14-DEC-1994;
                                                                                                                          04-MAY-1995;
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                 10-JUN-2003
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The present invention relates to a novel method for modulating growth, differentiation or survival of a cell. The method involves contacting the cell with a hedgehog polypeptide such as Desert hedgehog (Dhh), Sonic hedgehog (shh) and indian hedgehog (inh). The method is used to induce a cell to differentiate to a neuronal cell phenotype. It is used to modulate skeletogenesis. The method is used to treat a degenerative disorders of the nervous system such as neuronuscular, autonomic or central nervous system such as neuronuscular, autonomic or disease, amyotrophic lateral sclerosis, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, Huntington's disease, multiple sclerosis, Pick's disease, neuronal degeneration associated with a neuronal damage resulting from trauma and neuronal damage resulting from trauma and neuronal damage resulting from anoxia-ischaemia. The invention is also used for antisense gene therapy. The present sequence is human shu DNA amplifying PCR primer. This sequence is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hedgehog polypeptide; tissue array generation; tissue array maintainance; hedgehog; human; PCR; primer; ss.
                                                                                                                                                                                                                                               Modulating cell growth, differentiation or survival, for treating neurodegenerative diseases, such as Alzheimer's or Parkinson's disease, comprises contacting the cell with a hedgehog polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.6%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 3.8; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human sonic hedgehog primer seg id 43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24 ACCGAGGGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACCGAGGCTGGGACGAAGATGGC 24
                                                                                                                                                                                  Tabin CJ
                                                                                                                                                                                                                                                                                                                  Example 5; Page 49; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                              93US-00176427.
94US-00356060.
95US-00435093.
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94US-00356060.
95US-00435093.
95US-00460900.
                95US-00462386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-DEC-2000; 2000US-00736476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD71413 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                Ingham FW, Mcmahon AP,
                                                                                                              (INGH/) INGHAM P W. (MCMA/). MCMAHON A P. (TABI/) TABIN C J.
                                                                                                                                                                                                                WPI; 2003-803151/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2003190696-A1.
                                            30-DEC-1993;
14-DEC-1994;
04-MAY-1995;
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14-DEC-1994;
04-MAY-1995;
05-JUN-1995;
                35-JUN-1995;
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(HARD ) HARVARD COLLEGE.
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Marti-Gorostiza Tabin CJ, Bumcrot DA, Mcmahon AP, Ingham PW,

WPI; 2003-831623/77

New nucleic acid encoding a hedgehog polypeptide having an amino acid sequence identical or homologous to a vertebrate hedgehog protein, useful for generating or maintaining an array of different vertebrate tissue in vitro and in vivo.

Example 5; SEQ ID NO 43; 118pp; English.

The invention describes an isolated nucleic acid encoding a hedgehog polypeptide having an amino acid sequence identical or homologous to a vertebrate hedgehog protein or its portion and not identical to a fully defined 471-bp sequence. The nucleic acid is useful for generating and/or maintaining an array of different vertebrate tissue both in vitro and in vivo. This sequence represents a primer used to isolate DNA encoding vivo. This sequence r human sonic hedgehog.

Seguence 24 BP; 6 A; 5 C; 11 G; 2 T; 0 U; 0 Other;

Gaps ö 5.6%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 3.8; 7ative 0; Mismatches 0; Indels 24; Conservative Query Match Best Local Similarity Matches

## 47 24 ACCGAGGGCTGGGACGAAGATGGC

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24 ACCGAGGCTGGGACGAAGATGGC

RESULT

AAV18406 standard; cDNA; 25 AAV18406;

BP.

(first entry) 14-SEP-1998 Human mutated sonic hedgehog (SHH) gene exon 2 PCR primer.

Sonic hedgehog; SHH gene; HH gene; tumorigenesis; oncogenesis; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR; primer; ss 

Synthetic

Homo sapiens,

WO9821227-A1

22-MAY-1998

12-NOV-1997;

96US-00748591. 13-NOV-1996;

REGC ) UNIV CALIFORNIA.

WPI; 1998-297857/26.

Bonifas J;

Hu 2,

Epstein E,

New nucleic acid encoding oncogenic human hedgehog protein - usefue.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation.

Example; Page 23; 47pp; English.

used with This human sonic hedgehog (SHH) gene exon 2-specific primer was used v another exon 2-specific primer (see AAV18406) in a PCR using DNA from

human bacterial artificial chromosome (BAC) DNA pools. Only pools comprising a BAC that contains the sequence tag defined by the primer pair will yield an amplification product. The process was continued until a single positive BAC was identified. The positive clone, BAC270A17, was digested with restriction enzymes and ligated into vectorette linkers. Mutations (see AAV18403 and AAV18404) have been identified in the SHH gene in human cancers. The mutated SHH genes and the encoded polypeptides (see AAW48735 and AAW48735) can be used in methods for the treatment and diagnosts of cancer and other diseases involving cell proliferation or differentiation 8X3333333333X8

Sequence 25 BP; 4 A; 8 C; 8 G; 5 T; 0 U; 0 Other;

Gaps ô Score 23.4; DB 1; Length 25; Pred. No. 5.5; 0; Mismatches 1; Indels / Match Local Similarity 96.0%; les 24; Conservative Query Match Matches

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116 CAGCAAGTACGGCATGCTGGCCCGC 140 cascaastacsscarscrescresc 25

ð d RESULT 19

BP. ABZ79785 standard; DNA; 24 ABZ79785 ID ABZ7

ABZ79785;

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(first entry) 12-MAY-2003 Indian hedgehog PCR primer SEQ ID NO:5.

primer; Osteopathic; antirheumatic; antiarthritic; cytostatic; cartilage; cartilage differentiation; joint disease; bone fracture; myeloma; osteoporosis; rheumatoid arthritis; human; Indian hedgehog; PCR p

sapiens Synthetic. Omor

WO2003000870-A1.

03-JAN-2003

25-JUN-2002; 2002WO-JP006351

26-JUN-2001; 2001JP-00193503

(TAKE ) TAKEDA CHEM IND

Hikichi Y, Inazuka M;

WPI; 2003-201422/19

Culture method for cartilage differentiation from cells under hypoxic conditions into cartilage cells applicable in cartilage transplantation, and studying genes or proteins relating to joint diseases. 

Example 3; Page 29; 37pp; Japanese.

The present invention describes a method for cartilage differentiation by culturing cells capable of differentiating into cartilage under hypoxic conditions. Also described: (1) a method for producing cartilage cells or cartilage by culturing the required cells under hypoxic conditions; (2) drugs containing the produced cartilage cells or cartilage. (3) a method for preventing or treating joint diseases by transplanting an effective amount of the cartilage cells or cartilage; (4) the use of the cartilage cells or cartilage; (5) a method for screening genes relating to cartilage differentiation or joint diseases by using any of the culture methods; (6) a method for screening promoters or inhibitors of cartilage differentiation by using any of the culture methods; (7) a method for screening preventure methods; (7) a method for screening preventure methods; (7) a method for screening preventure methods; (8) a method for screening preventure methods; (9) a method for screening preventure methods; (1) a method for screening preventure methods; (2) a method for screening preventure methods; (3) a method for screening preventure methods; (1) a method for screening preventure methods; (2) a method for screening preventure methods; (3) a method for screening preventure methods; (2) a method for screening preventure methods; (3) a method for screening preventure methods; (2) a method for screening preventure methods; (3) and screening preventure methods; (3) a method for screening preventure methods; (4) a method for screening preventure methods; (5) a method for screening preventure methods; (6) a method for screening and screening preventure methods;

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Gaps

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Sequence 19 BP; 3 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

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Example; Page 23; 47pp; English.
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AAV18410 standard; cDNA; 19 BP.
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Best Local Similarity
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110 AAV10/

AAV10/
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cDNA derived from human epidermal keratinocytes was amplified by 3-stage nesting using sonic hedgehog (SHH) gene stage 1 primers (see AAV18413 and AAV18414).

AAV18414), stage 2 primers (see AAV18415 and AAV18416) and stage 3 primers (see AAV18415 and AAV18415). The PCR product was identified as authentic SHH, A single somatic mutation (see AAV18403) of the SHH gene was found in cancers arising from 3 different tissues in independent patients. Another mutation (see AAV18404) was identified in another cancer. The mutated SHH genes and the encoded polypeptides (see AAV48735) and AAV48736) can be used in methods for the treatment and diagnosis of cancer and other diseases involving cell proliferation or differentiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sonic hedgehog; SHH gene; HH gene; tumorigenesis; oncogenesis; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New mucleic acid encoding oncogenic human hedgehog protein - useful for, e.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation.
                                                                           4.5%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 23; Uive 0; Mismatches 0; Indels
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.00.0%; Pred. No. 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human mutated sonic hedgehog (SHH) gene PCR primer.
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                                                                                                                                                                               194 CCACTGCTCGGTGAAGCA 212
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                                                                                                                                                                                                                        CCACTGCTCGGTGAAAGCA 1
                                                                                                                                                                                                                                                                                                                                                   AAV18416 standard; cDNA; 19 BP.
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                                                                                                                                  Conservative
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                                                                         Query Match
Best Local Similarity
Matches 19; Conserval
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer; ss.
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Matches
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methods; (8) drugs containing the screened promoters or inhibitors of cartilage differentiation, or preventives or remedies for joint diseases; (9) a method for preventing or treating joint diseases by administering an effective dose of the promoters or inhibitors, or preventives or remedies to mammals; and (10) the use of the promoters or inhibitors, or preventives or remedies for producing drugs for joint diseases. The produced cultured cartilage cells or cartilage can be used in cartilage transplantation, studying genes or proteins relating to joint diseases and screening drugs for their treatment, including diseases of bone fracture, myeloma, osteoporosis and rheumatoid arthritis. The present sequence represents a PCR primer for Indian hedgehog, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sonic hedgehog; SHH gene; HH gene; tumorigenesis; oncogenesis; basal cell carcinoma; breast cancer; medulloblastoma; tumour; cell proliferation; cell differentiation; diagnosis; therapy; human; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid encoding oncogenic human hedgehog protein - useful for, e.g. treatment and diagnosis of cancer and diseases involving cell proliferation or differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 3 A; 4 C; 10 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                        Score 21.4; DE; Pred. No. 13; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              150 GAGGCCGCTTCGACTGGGTGTA 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAGGCCGGCTTTGACTGGGTGTA 23
                                                                                                                                                                                                                                                                                                                                                                             5.0%;
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Gaps

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284 CACCAAGCTGGTGAAGGAC 302

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CACCAAGCTGGTGAAGGAC

13

22 RESULT

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BST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human; cross-species comparison.
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                                                                                                                                                                                                                                                 16-MAR-2001; 2001US-0276759P.
                                                                                                                                                                                                             15-MAR-2002; 2002US-00098263.
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hes 21; Conservative
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                                                                                                                                  JS2003104410-A1.
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                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                           Mittmann MP;
                                                                                                                                                                      05-JUN-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is conceins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7922.1, MD24 is encoded at chromosome 7922.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15p11.2 and MD212 is encoded at chromosome 15p11.2 and MD212 is encoded at chromosome 15p11.2 and MD212 is encoded at chromosome or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23. MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for disgnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene expression. The proteins are useful as therapeutic agents for gene expression. The vaccines. The present sequence was used to illustrate the invention.
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                                                                                                                                                                      Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ12 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human microarray DNA oligonucleotide SEQ ID NO 66408.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 25 BP; 4 A; 11 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                    Human MDZ3 scanning oligonucleotide SEQ ID 1905.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          th 4.4%; Score 18.6; D
Similarity 84.0%; Pred. No. 53;
21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 8; SEQ ID NO 1905; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGTICCICACIAICCIGCCCGCGA 25
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                    BP.
                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                                                                                                                                                          32-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shannon M, Gu Y, Nguyen C;
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                    ADB00919 standard; DNA; 25
                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VPI; 2003-423107/40
                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                        3P1281758-A2
                                                                                             20-NOV-2003
                                                                                                                                                                                                                                                                                                                                               05-FEB-2003
                                                          ADB00919;
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Best Local
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ACI66417
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ADB00919
ID ADB0
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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its consequences including one of 2,018,500 fully defined sequences, or its consequence match, perfect mismatch, antisense match or antisense mismatch.

Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation to a DNA library, or at least one target sequence. The method of analysis comprises of at least one target sequence. The method of analysis comprises of hybridising at least one or more nucleic acids to at least two or more nucleic acids to probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms, or family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid further comprises a tag sequence. The array of nucleic acid further comprises a tag sequence. The array of nucleic acid further comprises a tag sequence. The array of nucleic acid further comprises a tag sequence or specific comparations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones of for additional subclones containing segments of DNA that have been contained and previously sequenced in the microarray. Note: The sequence of the form the form at a pardiar an also be obtained in electronic format directly from them instructions of the sequence of the format directly an also manience when the comparation is a series.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Desert hedgehog; human; HuDHH; PCR; RACE; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25 BP; 7 A; 8 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rom USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Desert hedgehog gene sense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.4%; Score 18.6; D
34.0%; Pred. No. 53;
tve 0; Mismatches
                                                                                                                                                                                                                             Claim 1; SEQ ID NO 66408; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      386 CGACGCCCAAGAAGGTCTTCTAC
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This sense primer corresponds to nucleotides 460-479 of a cDNA clone (see AAV62396) coding for novel human Desert hedgehog protein (see AAW79596).

I was used with an antisense primer (see AAV6211) in a first-steep PCR amplification of human leukaemia plasma cell line ARH-77 (ATCC CRL-1621) cDNA in a modified PCR method of 3'RACE. 2 Subsequent PCR amplifications (see AAV62423-26) yielded a cDNA clone (see AAV6239) encoding a C-terminal fragment (see AAW79599) of the novel human Desert hedgehog protein. Nucleotide sequences (see AAV62339-95) encoding mature and precursor forms (see AAW79593-55) of human Desert hedgehog are claimed. The Desert hedgehog DNA, protein and a claimed monoclonal antibody can be used in to elucidate hereditary morphological abnormalities in humans to establish their treatments and diagnoses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer, neuroectoderm cell; cell production; Parkinson's disease; early primitive ectoderm-like cell; EPL cell; cell therapy; transgenic animal; gene therapy; neuronal disease; Huntington's disease; lysosomal storage disease; multiple sclerosis; memory disorder; behavioural disorder; Alzheimer's disease; organ transplant;
                                                                                                                                                                                                                                                                                                     Human Desert hedgehog protein - and corresponding DNA and monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                  Example 1-4; Page 10; 39pp; English.
                                                                                                                                                                                                      (HAYB ) HAYASHIBARA SEIBUTSU KAGAKU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    156 GGCTTCGACTGGGTGTACTA 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GGCTTCGACTGGGTCTACTA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         behavioural disorder; Alzheim: spinal cord disorder; Shh; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF87046 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JAN-2001; 2001WO-AU000030.
                                                                                                                           98EP-00303187
                                                                                                                                                         97JP-00121578.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer for Shh gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                     WPI; 1998-544642/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200151611-A1.
                              Homo sapiens.
                                                                                                                                                         25-APR-1997;
14-APR-1998;
                                                                                                                           24-APR-1998;
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                                                          EP874048-A2.
                                                                                         28-OCT-1998
                                                                                                                                                                                                                                      Ariyasu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF87046;
               Synthetic
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                                                                                                                                                                                                                                                                                                                      antibody.
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AAF87046/
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This sequence represents a PCR primer for the Shh gene, used within the scope of the invention. The invention relates to a method for producing scope of the invention. The invention relates to a method for producing concrected early primitive ectoderm—like (EPL) cells and a neural-inducing conditioned primitive ectoderm—like (EPL) cells and a neural-inducing conditioned confident to a mine sufficient to generate controlled differentiation to (I). The cells or partially differentiated progeny are differentiation to animal cell therapy, transgenic animal production, confiderentiated progens in neuroectoderm cells or their partially differentiated progeny and evaluation of fearth of producing confiderentiated progeny and evaluation of pological molecules that direct differentiation of neural cells. The method is useful for producing concrectederm cells. It is also useful for producing differentiated cells from neural ectoderm cells. The method can consider the neuron of neuroectoderm cells in vitro in concrectederm cells from neuroectoderm cells in vitro in concrectederm cells from also be used for producing concreted disease, including Parkinson's disease, concretederm cells from the cells can be used in the concrete particular diseases, including Parkinson's disease, concretederm cells produced by the method are used for transplant. Neural concreted mentors and Schwann cells produced by the method are used for the transmit of confidered method are used for the transmit of confidered method are used for the method are used for the method are used for the particular of method are used for the transmit of method are used for the method ar
                                                                                                                                                                                                                            Producing neuroectoderm cells for treatment of Parkinson's and Alzheimer's and for transplantation comprises culturing early primitive ectoderm-like cells in conditioned medium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ds; Hedgehog protein; cancer; PCR; primer; amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hedgehog protein derivative primer 2.
                                                                                                                                                                                                                                                                                                                         Example 3; Page 41; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             355 ACAGCGACTICCICACTITC 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 acadedacricercaecric 1
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14-JAN-2000; 2000AU-00005098.
20-APR-2000; 2000AU-00007045.
27-APR-2000; 2000AU-00007143.
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                                                                                                                                    Rathjen J;
                                                                                          (BRES-) BRESAGEN LTD
                                                                                                                                                                                WPI; 2001-432908/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP10215867-A.
                      20-APR-2000;
27-APR-2000;
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                                                                                                                                    Rathjen PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-AUG-1998.
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Matches
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Gaps

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4.3%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 35; 1; Indels ive 0; Mismatches 1; Indels

Wed Apr

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encoding it - useful for prediction 9. lung cancer.

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Hedgehog protein derivative and gene en
and diagnosis of various diseases e.g.
                                                                                                                                                                                                      Disclosure; Page 6; 7pp; Japanese
(ASAG ) ASAHI GLASS
                                                     WPI; 1998-499061/43.
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The primers AAV59457-V59462 were used in the production of hedgehog (hh) protein derivative may be used in the prediction and diagnosis various diseases e.g. cancer Length 25; 3; Indels Sequence 25 BP; 3 A; 9 C; 8 G; 5 T; 0 U; 0 Other; DB 1; Score 18.2; DE Pred. No. 64; 0; Mismatches 244 222 GIGGCGGCCAAATCGGGAGGCTG Match 4.3%; Local-Similarity 87.0%; hes 20; Conservative Query Match Best Local-Si Matches 20

24 gredecedecharicegaedecre 2 à

ADB00921 standard; DNA; 25 BP. 20-NOV-2003 Human MDZ3 ADB00921; RESULT 27 4DB00921 

scanning oligonucleotide SEQ ID 1907. (first entry)

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens

SP1281758-A2

05-FEB-2003,

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC

ΰ Nguyen Gu ₹, Shannon M,

WPI; 2003-423107/40.

MDZ3, New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1907; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7g22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 15g26.1. The MD23, MD24, MD27, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic

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                                                   probes are
acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes ar useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                               Query Match
4.3%; Score 18.2; DB 1; Length 25;
Best Local Similarity 87.0%; Pred. No. 64;
Matches 20; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                    Sequence 25 BP; 3 A; 11 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  363 TICCICACITICCIGGACCGCGA 385
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BP. 25 (first entry) ADB00920 standard; DNA; 20-NOV-2003 ADB00920; RESULT 28 ADB00920

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Gaps

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Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Human MDZ3 scanning oligonucleotide SEQ ID 1906.

Homo sapiens 

EP1281758-A2 05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181. (AEOM-) AEOMICA INC

Gu Y, Nguyen C; Shannon M,

WPI; 2003-423107/40.

MDZ3 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1906; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome FQ22.1, MD24 is encoded at chromosome FQ22.1, MD24 is encoded at chromosome 16pl.1.2 and MD212 is encoded at chromosome 16pl.1.2 and MD212 is encoded at chromosome 16pl.1.2 and MD212 is encoded at chromosome 16pl.2 sequences are useful in therapy, continuated with decreased or increased expression or archivity of MD23, associated with decreased or increased expression or activity of MD23, continuated with decreased or increased expression or activity of MD23, consect or developmental disorders. The nucleic caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The nucleic useful in constructing microarrays for measuring gene expression. The proteins are useful as therapoutic agents for gene therapy or as proteins are useful as therapoutic agents for gene therapy or as

BP; 3 A; 11 C; 5 G; 6 T; 0 U; 0 Other; Sequence 25 ADD15351;

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Indels

Length 25;

DB 1;

4.3%; Score 18.2; Local Similarity 87.0%; Pred. No. 64; es 20; Conservative 0; Mismatches

Query Match

Matches

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AAH45474 standard; DNA; 18 BP.

AAH45474/c

RESULT 29

(first entry)

07-SEP-2001

AAH45474;

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This invention relates to a novel method for the detection, treatment and/ or prevention of callular debilitations or derangements caused by the development of sporadic basal cell carcinoma (BCC). Specifically, it refers to the identification of relevant theirapeutic agents based on their effect on the expression level and activity of the Glil or transsription factor gene. Glil is a proto-oncogene that is ectopically expressed in epidermal tissue and is linked to tumour formation and neoplasia. The present invention describes cytostatic Glil inhibitors that are useful for detecting the onset or presence of sporadic BCC in an animal. Furthermore, it includes methods for testing the ability of a nituy or other entity to endulate the activity of Glil. This oligonucleotide sequence is the RT-PCR primer Shh-D used to amplify human Shh (secreted sonic hedgehog) RNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Upstream activating sequence; transgenic animal; regulatory DNA sequence; hedgehog gene; bigenic animal; transcriptional activating sequence; disease model; cancer; altered vascularisation; brain size regulation; autoimmune disease; tissue proliferation; Parkinson's disease; Shh;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preventing or treating sporadic basal cell carcinoma by administering an inhibitor of glioma transcription factor-1 (Gli1) activity or expression, and diagnosis of the disease by detecting the presence and level of expression of Gli1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                 RT-PCR; primer; Shh-D; human; ss; PCR; cellular debilitation; sporadic basal cell carcinoma; BCC; Glil; proto-oncogene; tumour formation; neoplasia; cytostatic; secreted sonic hedgehog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 33; 0; Indels ive 0; Mismatches 0; Indels
                                                                                           RT-PCR primer Shh-D used to amplify human Shh RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 4 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 6; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 deadricricrecactacea 1
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                                                                                                                                                                                                                                                                                                                                                                                                                          97US-0050286P.
98US-00102491.
                                                                                                                                                                                                                                                                                                                                                                               03-APR-2001; 2001US-00825155.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  59 GGAGTCTCTGCACTACGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                111/c
AAZ49111 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                             (first entry)
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Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ALTA/) ALTABA A R I.
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                                                                                                                                                                                                                                                                                    US2003100032-A1.
                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                          20-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-APR-2000
                                           15-JAN-2004
                                                                                                                                                                                                                                                                                                                               29-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a method of detecting the onset or presence of sporadic basal cell carcinoma (BCC) in an animal. The method involves measuring the level of Gili in a sample of Skin. Gili levels above basal or normal indicate the presence or onset of sporadic basal cell carcinoma. Gili is a zinc finger transcription factor down stream of secreted sonic hedgehog (shh) activation in a cascade of cytoplasmic signal transduction. Gili in turn can induce Shh expression in an autoregulatory manner. There are links between ectopic expression of the Gili gene and the development or onset of BCC. The method is useful for detecting the onset or presence of sporadic basal cell carcinoma, particularly in detecting skin cancer. The present sequence represents a PCR primer specific for human Shh cDNA. The primer is used in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting the onset or presence of skin cancer, particularly sporadic basal cell carcinoma, comprises measuring the level of Glil in the
                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sporadic basal cell carcinoma, BCC, detection, Glil, skin cancer, transcription factor, PCR primer, human, 88; sonic hedgehog; shh.
                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer Shh-D specific for human secreted sonic hedgehog cDNA.
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Disclosure; Col 8; 21pp; English.

sample.

98US-00102491.

22-JUN-1998;

29-MAY-2001

20-JUN-1997;

Homo sapiens,

US6238876-B1

(UYNY ) UNIV NEW YORK STATE.

WPI; 2001-366473/38.

Altaba ARI;

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4.2%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 33; tive 0; Mismatches 0; Indels

16 Н

Conservative

Similarity

Query Match Best Local Simi Matches 18;

GGAGTCTCTGCACTACGA GGAGTCTCTGCACTACGA

13

8

BP.

ADD15351/c ID ADD15351 standard; DNA; 18 XX

RESULT 30

Sequence 18 BP; 4 A; 5 C; 5 G; 4 T; 0 U; 0 Other;

the invention

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This sequence represents a PCR primer for the mouse Shh gene. The invention relates to a transgenic non-human animal (A) whose cells contain a non-viral regulatory DNA sequence (I) (e.g. an upstream activating sequence) linked to a recombinant hedgehog gene (II), which was introduced into the mammal, or its ancestor, at an embryonic stage. Bigenic animals (A'), derived from (A) by introducing a transcriptional activating sequence (TAS), are useful as models of disease, particularly cancer (of breast, skin, prostate, kidney, lung, or central nervous system, also primitive neuroectodermal tumours and medulloblastomal. CC cancer (of breast, skin, prostate, kidney, lung, or central nervous crarget genes on signaling pathways involving hedgehog proteins (HP) (e.g. altered vascularisation, regulation of brain size, density and cellular pathways involving hedgehog proteins (HP) (e.g. altered vascularisation, regulation of brain size, density and cellular pathways involving for a temporal requirement for HP in disease progression (particularly of cancers and autoimmune disease). The animals can be used to screen for potential therapeutic proliferation and differentiation. Hedgehog proteins can also be used to expand a population of neural stem cells from a subject, then the cells consistent of the subject, specifically for treatment of Parkinson sor and the silent transgenes in progeny from a simple cross since the transcription activator and the silent transgene are maintained in separate mouse lines, and abnormal expression consistent embryon and genotypic screening for each experiment, and many microin embryon and genotypic screening for each experiment, and many microin embryons and many procession experiment.
Alzheimer's disease; spinal cord injury; therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                Transgenic animals useful as disease models, e.g. for cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oigenic embryos can be produced by cross-breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 5 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 20; 44pp; English.
                                                                                                                                                                                        99WO-US012417.
                                                                                                                                                                                                                                        98US-0087899P.
                                                                                                                                                                                                                                                                                                                                    Rowitch DH, Mcmahon AP;
                                                                                                                                                                                                                                                                                    HARD ) HARVARD COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-105693/09.
                                                                                            WO9963052-A2
                                                                                                                                                                                          1666T-NDD-E0
                                                                                                                                                                                                                                        03-JUN-1998;
                                                                                                                                          39-DEC-1999
                                                Mus sp.
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The present invention describes the rat Nurrl coding and protein sequences. The Nurrl protein is involved in the induction of tyrosine Approxylase expression in adult rat-derived hippocampal progenitor cells. The Nurrl gene and protein can be used in the treatment of catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia. They can also be used to induce tyrosine hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor cells

Sequence 21 BP; 2 A; 7 C; 5 G; 7 T; 0 U; 0 Other;

Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia.

Example 1; Page 20; 68pp; English

(SALK ) SALK INST BIOLOGICAL STUDIES.

Palmer T,

Sakurada K,

WPI; 2000-656165/63.

21-MAR-2000; 2000WO-US007544

05-OCT-2000

26-MAR-1999;

Rattus norvegicus. WO200058451-A1.

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Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD27; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                        Gaps
                        ö
4.2%; Score 17.8; DB 1; Length 21; 90.5%; Pred. No. 52; tive 0; Mismatches 2; Indels
                                                                                                                                                                                             Human MDZ3 scanning oligonucleotide SEQ ID 1904.
                                             104 TGACCGCGACCGCAGTA 124
                                                                    TGACAGGGACCGCAGCAAGTA 1
                                                                                                                            ВÞ.
                                                                                                                                                                                                                                                                                                                                             30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                                                                                                  02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                               Gu Y, Nguyen C;
                                                                                                                             ADB00918 standard; DNA; 25
                                                                                                                                                                        (first entry)
                          19; Conservative
  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-423107/40.
                                                                                                                                                                                                                                                                                                                                                                                          (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                  EP1281758-A2.
                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                        20-NOV-2003
                                                                                                                                                                                                                                                                                                                       05-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                               Shannon M,
                                                                                                                                                  ADB00918;
                                                                    21
                          Matches
                                                                                                      RESULT 33
                                                                                                                  ADB00918
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Gaps

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DB 1; Length 21; 2; Indels

Pred. No. 52; 0; Mismatches 4.2%; Score 17.8; F

Local Similarity 90.5%; les 19; Conservative

Query Match

Best Loc Matches

57 GAGGAGTCTCTGCACTACGAG 77

8

21 GAGGAGTCTCTACACTATGAG 1

ВЪ

AAA95383 standard; DNA; 21

RESULT 32 AAA95383, Rat; Nurrl; tyrosine hydroxylase; catecholamine-related disease; Parkinson's disease; manic depression; schizophrenia; PCR primer; ss.

Rat Shh coding sequence PCR primer #2.

(first entry)

12-FEB-2001

AAA95383;

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New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1904; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is concoded at chromosome fg21.2, MD27 is encoded at chromosome fg21.2, and MD212 is encoded at chromosome is foll.2 and MD212 is encoded at chromosome is foll.2 and MD212 is encoded at chromosome is 5g26.1. The MD23, MD24, MD27, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23 ymD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are protein are useful as therapeutic agents for gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.

Sequence 25 BP; 3 A; 12 C; 4 G; 6 T; 0 U; 0 Other;

Gaps .; 0 DB 1; Length 25; 4; Indels 4.1%; Score 17.6; D 83.3%; Pred. No. 85; :ive 0; Mismatches 361 ACTICCICACITICCIGGACCGCG 384 2 AGTICCICACIAICCIGCCCGCG 25 Query Match
Best Local Similarity 83.3
Matches 20; Conservative ò 셤

RESULT 34 ABS55991/c ID ABS55991 standard; DNA; 22 BP.

ABS55991; THE STANK BY SECTION S

23-JAN-2003 (first entry)

Mouse RT-PCR primer Shh rp #1.

Mouse; primer, ss; Hedgehog signalling pathway; T-cell mediated disease; T-cell apoptosis; Notch signalling pathway; cancer; breast; prostate; ovary; T-cell activation; T-cell proliferation; lymphoma; carcinoma; autoimmune disease; inflammatory disease; proliferative disorder; viral infection; genetic immunodeficiency; neurodegenerative disease; myelodysplastic syndrome; isohemic injury; toxin-induced disease; wasting disease; RT-PCR; reverse transcriptase; Shh; sonic hedgehog.

Mus musculus

WO200280952-A2

09-APR-2002; 2002WO-GB001666.

09-APR-2001; 2001GB-00008872. 09-APR-2001; 2001GB-00008873.

(LORA-) LORANTIS LTD.

Champion BR; Lamb JR, Hoyne GF, Dallman MJ,

WPI; 2003-058470/05

Use of a modulator of Hedgehog signaling pathways for treating T-cell mediated disease or infection and diseases associated with increased or decreased T-cell apoptosis and T-cell proliferation.

The invention relates to use of a modulator of a Hedgehog signalling pathway or a modulator of a target of the pathway in the preparation of a medicament for treating T-cell mediated disease or infection or a disease or disorder associated with increased or decreased T-cell apoptosis and for modulation or the Notch signalling pathway in cell apoptosis, and for modulation of the Notch signalling pathway in cimume cells. The modulator is useful for treating cancer of the breat, prostate or ovary, lymphomas and carcinomas, autoimmune diseases such as systemic lupus erythematosus, multiple sclerosis and diseases such as opposite or proliferative disorders such as obtending and crohn's disease, inflerative disorders such as atherosolerosis and psoriasis, viral infections such as AlDS and herpesviruses, genetic immunodeficiencies, carcodegenerative diseases such as Alzheimer's disease and Parkinson's disease, myelodysplastic syndromes such as aplastic anaemia, isochamic cirrhosis and wasting diseases such as plastic induced diseases such as carcheria this sequence represents increased a reverse transcriptase PCR (RT-PCR) primer used in the scope of the Example 10; Page 110; 154pp; English. 

Sequence 22 BP; 6 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

Gaps ö DB 1; Length 22; 3; Indels 4.0%; Score 17.2; L 86.4%; Pred. No. 76; tive 0; Mismatches Best Local Similarity 86.4 Matches 19; Conservative Query Match

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RESULT 35

ADB00922 standard; DNA; 25 ADB00922

ADB00922;

(first entry) 20-NOV-2003 Human MDZ3 scanning oligonucleotide SEQ ID 1908.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD212; Chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874. 

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

ີ Shannon M, Gu Y, Nguyen

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 1908; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is

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encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids and proteins are also beful for diagnosing or monitoring a disease acids and proteins are also beful for diagnosing or monitoring a disease acids and bo be used as probes to detect and characterize gross alterations in MDZ3, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.

Sequence 25 BP; 3 A; 11 C; 4 G; 7 T; 0 U; 0 Other;

Query Match
4.0%; Score 17.2; DB 1; Length 25;
Best Local Similarity 66.4%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 3; Indels 364 TCCTCACTTTCCTGGACCGCGA 385 ò

1 rccrcacrarccraccccccc a

ACK14726 standard; DNA; 25 BP. RESULT 36 ACK14726

14-OCT-2003 (first entry) ACK14726;

Human microarray DNA oligonucleotide SEQ ID NO 114707.

expressed sequence tag; microarray; gene expression; on; biallelic marker; polymorphism; human; cross-species comparison. EST; ss; probe; exg genetic variation;

Homo sapiens.

05-JUN-2003.

US2003104410-A1.

15-MAR-2002; 2002US-00098263

16-MAR-2001; 2001US-0276759P.

(AFFY-) AFFYMETRIX INC.

Mittmann MP;

WPI; 2003-567953/53.

New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.

Claim 1; SEQ ID NO 114707; 9pp; English.

The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect manch, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises hybridising at least one or more nucleic acids to at least two or more nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms,

or family members of a gene and a cross-species comparison. Bach of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotblot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening CDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly from USPTO at sequence.html

88998888888888888

Sequence 25 BP; 5 A; 9 C; 6 G; 5 T; 0 U; 0 Other;

ö 4.0%; Score 17.2; DB 1; Length 25; 86.4%; Pred. No. 1e+02; 3; Indels 0; Mismatches 19; Conservative Similarity Query Match Best Local S Matches

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à

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Gaps

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RESULT 37 AAA15463

BP. AAA15463 standard; DNA; 25

AAA15463;

(first entry) 21-SEP-2000 PCR primer for a rat connective tissue growth factor DNA.

Rat, connective tissue growth factor; CTGF; cell proliferative disorder; connective tissue cell; scleroderma; arthritis; cirrhosis; hepatic fibrosis; renal fibrosis; atherosclerosis; cardiac fibrosis; adhesion; surgical scarring; PCR primer; ss.

Rattus sp.

WO200027868-A2.

18-MAY-2000.

99WO-US026189. 05-NOV-1999;

98US-00187478. 99US-00292036. 06-NOV-1998; 14-APR-1999;

(FIBR-) FIBROGEN INC.

Sverdrup F, Carmichael DF; Schmidt BF, Allen ML,

WPI; 2000-376484/32.

New rat connective tissue growth factor, its related gene and antisense sequences useful for modulating CTGF and treatment of cell proliferative disorders. 

Example 1; Page 37; 55pp; English.

PCR primers AAA15463-64 were used to amplify DNA encoding a rat connective tissue growth factor (CTGF) polypeptide. The polypeptide may blay a significant role in the normal development, growth and repair of mammalian tissue. Antisense sequences can be used to inhibit the expression of CTGF in a cell. In particular, the antisense sequences are useful for ameliorating cell proliferative disorders associated with CTGF, e.g. overgrowth of cells, e.g. connective tissue cells. The regulation of CTGF activity comprises down-regulation. The disorders, which can be treated, are chosen from soleroderma, arthritis, cirrhosis, hepatic fibrosis, renal fibrosis, atherosclerosis, cardiac fibrosis, adhesions and surgical scarring. The amisense sequences can also be used to detect expression of CTGF in a sample

BST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human; cross-species comparison.

US2003104410-A1. Homo sapiens.

05-JUN-2003.

Human microarray DNA oligonucleotide SEQ ID NO 66407.

(first entry)

14-OCT-2003

ACI66416;

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ACI66416 standard; DNA; 25 BP.

RESULT 39 ACI66416

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PCR primers AAZ99759-60 were used to amplify a 558 bp of the connective tissue growth factor (CTGF) gene. The specification describes methods for treating or preventing fibrosis or a renal disorder associated with overproduction of extracellular matrix, by administering to a subject an agent that modulates, regulates, or inhibits the expression or activity of CTGF, Healthy individuals demonstrate consistently low levels of urinary CTGF, while in patients with kidney disease the mean level of CTGF increased 4-fold. In those patients with diabetes, but as yet undiagnosed kidney disease, a similar increase was seen. The methods and agents are useful for diagnosing, treating or preventing fibrosis, diabetes, hypertension or a renal disorder associated with overproduction of extracellular matrix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Diagnosing, treating or preventing fibrosis, diabetes or a renal disorder associated overproduction of extracellular matrix comprises administering an agent which modulates/inhibits the expression/activity of connective
                                                                                                                                                                                                                                                                                                                                                                                 Connective tissue growth factor; CTGF; fibrosis; renal disorder; extracellular matrix; kidney disease; diabetes; hypertension; FCR primer;
                                                                                                                                                                                                                                                                                                                                                   primer F used to amplify a 558 bp fragment of the CTGF gene.
                                                                                     ;
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Pred. No. 1.1e+02;
0; Mismatches 5; Indels
                                                  Length 25;
                                                                                   5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 BP; 5 A; 4 C; 12 G; 4 T; 0 U; 0 Other;
                Seguence 25 BP; 5 A; 4 C; 12 G; 4 T; 0 U; 0 Other;
                                              4.0%; Score 17; DB 1; 180.0%; Pred. No. 1.1e+02; ive 0; Mismatches 5,
                                                                                                                      162 GACTGGGTGTACTACGAGTCCAAGG 186
                                                                                                                                                      1 cagredererereaceaeccaaes 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 44; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (FIBR-) FIBROGEN INC.
(FORD-) FORD HEALTH SYSTEM HENRY.
                                                                                                                                                                                                                                             AAZ99759 standard; DNA; 25 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0099471P.
98US-0112855P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-US020601.
                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Riser BL, Denichilo M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tissue growth factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-256864/22.
                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200013706-A1.
                                                                                                                                                                                                                                                                                                                  12-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-SEP-1998;
16-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-MAR-2000.
                                                                                                                                                                                                                                                                                 AAZ99759;
                                                Query Match
Best Local (
                                                                    Best Loca
Matches
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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its correct match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in analysis of genetic variation or in hybridisation to a DNA library, and an monitoring gene expression levels by hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis compounds. The nucleic acid probes are specifically designed for marked compounds. The nucleic acid probes and detecting the hybridisation. The nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring compenses are attached to a solid support. The analysis comprises monitoring compenses are attached to a solid support. The analysis comprises monitoring can eamily members of a gene and a crose species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes in mapping the 5' termini of mRNA molecules by complete acide any gene, in mapping the 5' termini of mRNA molecules by conference and previously sequenced. The sequence or subclones containing segments of DNA that have been contained and previously sequenced. The sequence presented is one of the concleic acid probes incorporated in the microarray. Note: The sequence cata for this patent can also be obtained in electronic format directly concleic acid probes incorporated in the microarray. Note: The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25 BP; 7 A; 9 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 66407; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         386 cgaccaccaacaacarcrictac 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CGACGACCAACTAGGTCTTCGAC 25
                                                                                                                                                                                                                                                                                                                                                                            15-MAR-2002; 2002US-00098263.
                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAR-2001; 2001US-0276759P.
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Best Local Similarity 80.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-567953/53.
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Gaps

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162 GACTGGGTGTACTACGAGTCCAAGG 186

ઠે g

ch 4.0%; 1 Similarity 80.0%; 20; Conservative

Query Match Best Local Similarity Matches 20; Conserv

1 GAGTGGTGTGACGAGCCCAAGG 25

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New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.
                                                                                  expressed sequence tag, microarray; gene expression;
on; biallelic marker; polymorphism; human;
                                                                Human microarray DNA oligonucleotide SEQ ID NO 8430
                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 8430; 9pp; English.
              ACI08439 standard; DNA; 25 BP.
                                                                                                                                                                                       16-MAR-2001; 2001US-0276759P.
                                                                                                                                                                      15-MAR-2002; 2002US-00098263
                                                13-OCT-2003 (first entry)
                                                                                  EST, ss, probe, expressed
genetic variation, bialle
cross-species comparison.
                                                                                                                                                                                                       (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                          WPI; 2003-567953/53.
                                                                                                                                    US2003104410-A1.
                                                                                                                     Homo sapiens.
                                                                                                                                                      05-JUN-2003.
                                                                                                                                                                                                                         Mittmann MP;
                                ACI08439;
RESULT 40
        ACI08439
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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in analysis of genetic variation or in bybridisation to a DNA library, compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises of at least one or more nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring of mucleic acid probes and a gene and a gross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes in situ hybridisation, in Southern, Northern or dotablot hybridisation to identify or detect the sequence or specific or additional subclones containing segments of DNA that have been for additional subclones containing segments of DNA that have been contained and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly containing the formal previously sequence. The first containing the formal previously sequence them.
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Gaps
                                     ..
0
4.0%; Score 17; DB 1; Length 25;
80.0%; Pred. No. 1.1e+02;
live 0; Mismatches 5; Indels
                   l Similarity 80.0
20; Conservative
     Query Match
Best Local
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GACCGCGACGACGCCCAAGAAGG 402 GACCCCGACGTCGTCGCTAAGAGGG 25 378

à 셤 RESULT 41 AAF27037

멺. AAF27037 standard; DNA; 37

AAF27037;

(first entry) 30-MAR-2001 Human Sonic hedgehog (Shh) mutagenic primer, SEQ ID NO:41.

Sonic hedgehog; Shh; polymer conjugate; polyalkene glycol group; bloavailbhilty; formulation; neurological disorder; inflammarcry disorder; autoimmune disorder; cancer; neurodegenerative disorder; Parkinson's disease; Huntington's disease. Alzhedmer's disease; heurological injury; stroke; multiple sclerosis; multipnant glioma; medulloblastoma; neuroectodermal tumour; ss. mutagenic primer; ss. 

Homo sapiens Synthetic.

WO200073337-A1.

07-DEC-2000.

26-MAY-2000; 2000WO-US014741

99US-0137011P. 99US-0149016P. 01-JUN-1999; 13-AUG-1999;

(BIOJ ) BIOGEN INC.

Garber Taylor F, Pepinsky RB,

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WPI; 2001-049927/06

Modified hedgehog protein, useful in the treatment of Parkinson's disease and Huntington's chorea, comprises a polymer containing a polyalkylene glycol group linked to any residue other than the N-terminal and lysine residues.

Example 6; Page 77; 157pp; English.

increased thermal stability would be an advantage when formulating the hedgehog protein in powder form. The present sequence represents a human Sonic hedgehog mutagenic primer used in an exemplification of the protein of the invention to cross the blood-brain barrier, and an nvention

Sequence 37 BP; 6 A; 10 C; 12 G; 9 T; 0 U; 0 Other;

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ1; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

30-JUL-2002; 2002EP-00016874. 02-AUG-2001; 2001US-00922181

05-FEB-2003.

Homo sapiens. EP1281758-A2. Nguyen C;

Gu Y,

Shannon M,

WPI; 2003-423107/40.

(AEOM-) AEOMICA INC.

Human MDZ3 scanning oligonucleotide SEQ ID 1903.

(first entry)

20-NOV-2003

ADB00917;

BP.

ADB00917 standard; DNA;

ò

RESULT 43

vaccine; human;

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The invention relates to the use of an active agent stimulating the expression and/or function of serum response factor (SRP). SRP variants and/or members of the SRP signal transduction pathway in eukaryotic cells for the preparation of a therapeutic drug or a pharmaceutical composition for the treatment of disturbances or illness such as tumour invasion, tumour metastasis, auto-immune diseases, disturbances of would healing, lymphocyte homing and disturbances of immune defense mechanisms that are linked with SRP- related cellular malfunctions. Pharmaceutical compositions of the invention are used in treating diseases associated with expression or misexpression of SRP target gene, which include formation of diseases like metastatic cancer which is influenced by the gene uph-R, diseases like chronic renal failure, cancer and various hypoglycaemias. The present sequence is Shh specific reverse transcription PCR (RT-PCR) primer used in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of active agent stimulating expression of serum response factor, its variants or components of signal transduction pathway of factor in eukaryotic cells, for treating disturbances or illness e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                         tumour invasion; tumour metastasis; auto-immune disease; would healing; lymphocyte homing; immune defense mechanism; chronic renal failure; cellular malfunction; metastatic cancer; illness; hypoglycaemia; RT-PCR, Shh; reverse transcription PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                           Serum response factor; SRF modulator; signal transduction; disturbance;
                                         ;
0
    Length 37;
                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 23 BP; 4 A; 3 C; 8 G; 8 T; 0 U; 0 Other;
    4.0%; Score 17; DB 1; Le
69.7%; Pred. No. 2.6e+02;
iive 0; Mismatches 10;
                                                                              TGCTGGCCCGCCTGGCGGTGGAGGCCGGCTTCG 162
                                                                                                                   recadadacrecrecadregregecearerres
                                                                                                                                                                                                                                                                                                                                 specific reverse RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 7; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-SEP-2000; 2000EP-00119741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-SEP-2000; 2000EP-00119741
                                                                                                                                                                                               AAD34565/c
ID AAD34565 standard; DNA; 23
                                                                                                                                                                                                                                                                                            (first entry)
                      Similarity 69.7
3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NORD/) NORDHEIM A.
                                                                                                                                                                                                                                                                                            16-JUL-2002
Query Match
Best Local Simi
Matches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1186319-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nordheim A;
                                                                              130
                                                                                                                   'n
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                                                                                                                                                                             RESULT 42
                                                                                                                                                                                                                                                                                                                                   Shh
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD24, MD24, MD212, MD212. MD23 is cancded at chromosome 7422.1, MD24 is encoded at chromosome 6721.2, MD27 is encoded at chromosome 6721.2, MD27 is encoded at chromosome 16011.2 and MD212 is encoded at chromosome 16011.2 and MD212 is encoded at chromosome 5921.3-22.2, MD27 is encoded at chromosome 16011.2 and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder sessociated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 25 BP; 3 A; 12 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 8; SEQ ID NO 1903; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        361 ACTICCICACTITCCIGGACGGC 383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Agriccicacianiccioccocos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 82.6%;
les 19; Conservative
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Gape

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Score 16.6; DB 1; Length 23; Pred. No. 1.1e+02; 0; Mismatches 4; Indels

3.9%; 82.6%;

Query Match
Best Local Similarity 82.6
Matches 19; Conservative

177 GAGTCCAAGGCACATATCCACTG 199

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GAATCCAAAGCTCACATCCACTG 1

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The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch.

Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, or analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises of at least one target sequence. The method of analysis comprises of hybridisation are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis comprises monitoring probes are attached to a solid support. The analysis of nucleic acid for family members of a gene and a cross-species comparison. Each of the nucleic acide further comprises a tag sequence. The array of nucleic acide further comprises a tag sequence. The array of nucleic acide further comprises a tag sequence. The array of nucleic acide further comprises a tag sequence or specific.

Diot hybridisation to identify or detect the sequence or specific or additional subclones on in screening cDNA or genomic libraries or subclones for additional subclones containing sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence attached and previously sequenced in the microarray. Note: The sequence is received and compared to the microarray. Note: The sequence attached and previously sequenced in the microarray. Note: The sequence attached and sequence or also be obtained in electronic format directly and sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New array of nucleic acid probes, useful for in situ hybridization, in Southern, Northern or dot-blot hybridization to identify or detect the sequence or specific mutations of any gene.
                                                                   EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human; cross-species comparison.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
3.9%; Score 16.6; DB 1; Length 25;
Best Local Similarity 82.6%; Pred. No. 1.3e+02;
Matches 19; Conservative 0; Mismatches 4; Indels
                       Human microarray DNA oligonucleotide SEQ ID NO 14720.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25 BP; 5 A; 9 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 14720; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           251 GGGCTCGGCCACGGTGCACCTGG 273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gercreeceacearaaceree 1
                                                                                                                                                                                                                                                                                                             .5-MAR-2002; 2002US-00098263.
                                                                                                                                                                                                                                                                                                                                                            6-MAR-2001; 2001US-0276759P
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                                                                                                                                                                                                                                                                                                                                                                                                             (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-567953/53.
                                                                                                                                                                                                                US2003104410-A1
                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mittmann MP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-OCT-2003
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셤
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The present invention relates to nucleic acid sequences that are complementary to particular genes, and can be used as probes for a variety of analyses such as gene expression analysis. Each probe comprises 9 or more consecutive nuclectides from at least one of 14936 nuclectide sequences defined in the patent, or their perfect sense match, sense mismatch, antisense match or antisense mismatch oligonuclectides. The probes may be used in a array comprising at least 10 distinct nucleic acid probes. The array is useful in monitoring gene expression nucleic acid probes. The array is useful in monitoring gene expression the probes and in hybridisation to a DNA library, in analysing genetic cuseful for identifying family members of a gene. The probes are also useful in in situ hybridisations, in screening cDNA or genomic libraries (or derived subclones) for additional clones containing segments of DNA that have been previously isolated and sequenced, in Southern, northern, or dot-blot hybridisation of genomic DNA to identify or detect the sequence of any gene or detect specific mutations in any gene, and in mapping the 5' termin of many molecules by primer sextensions. The sequences of the invention are also useful as PCR primer complementary to particular genes with a wide range of analytical uses. Complemented for this patent was obtained in electronic format the sequence data for this patent was obtained in electronic format directly from the USPITO web site at sequences of the invention. Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New probe array useful e.g. for monitoring gene expression levels, for analyzing genetic variations, or for hybridizing tag-labeled compounds, comprises multiple nucleic acid probes.
               Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; library screening; Southern hybridisation; northern hybridisation; dot-blot hybridisation; gene sequence; mtation detection; target sequence; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer P24 to convert human antibody CAT-212 to IgG format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.9%; Score 16.6; DB 1; Length 25; 82.6%; Pred. No. 1.3e+02; Live 0; Mismatches 4; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 2 A; 9 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 2490; 9pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               167 GGTGTACTACGAGTCCAAGGCAC 189
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                                                                                                                                                                                                                                                                          08-AUG-2002; 2002US-00215112
                                                                                                                                                                                                                                                                                                                   08-AUG-2001; 2001US-0311040P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAD18152 standard; DNA; 21
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-576608/54
                                                                                                                                                                                                                                                                                                                                                            (MITT/) MITTMANN M.
                                                                                                                                                                                       US2003082596-A1
                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-DEC-2001
                                                                                                                                                                                                                                  01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                      Mittmann M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD18152;
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DNA target sequence #2490 useful in array for genetic analyses

The invention relates to a specific binding member which binds to human ectaxin. The binding member comprises an antibody variable heavy (VI) domain from CAT-212 (WIV) domain and a VH/VL (OMB) Variable light (VI) domain from CAT-212 (WIV) domain and a VH/VL (CDRs). Ectaxin is a chemoattractant protein that binds to a specific receptor which is expressed predominantly on eosinophils. The binding member is useful for neutralising ectaxin, which is useful in treating asthma, eczema and other atopic diseases such as rhinitis, food allergy, conjunctivitis, allergic colitis which are recognised as eosinophil-mediated diseases, for treating skin and other atopic conditions such as poriasis, pemphigoid, wells' syndrome, cellulitis, drug eruptions; inflammatory bowel disease which includes eosinophilic colitis/enteritis/ syndrome. The present sequence is a PCR primer used for converting encoding human antibody CAT-212 (ScPv-single chain variable region fragment) to IgG DNA (whole antibody) format Human, eotaxin; CAT-212; antibody; heavy chain variable region; VH; eczema; asthma; atopic disease; dermatological; rninitis; food allergy; vasotropic; conjunctivitis; allergic colitis; psoriasis; pemphigoid; eosinophil-mediated disease; cellulitis; drug eruption; vasculitis; inflammatory bowel disease; gastroenteritis; PCR primer; se. Human antibodies against ectaxin useful for treating asthma, eczema and other atopic diseases, comprises an antibody variable heavy or variable light domain from CAT-212 or from complementary determining regions. Sequence 21 BP; 3 A; 4 C; 11 G; 3 T; 0 U; 0 Other; (CAMB-) CAMBRIDGE ANTIBODY TECHNOLOGY Example 11; Page 103; 107pp; English Smith S; 02-MAR-2001; 2001WO-GB000927. 03-MAR-2000; 2000US-0187246P. Vaughan TJ, Wilton AJ, WPI; 2001-589944/66. WO200166754-A1. Homo sapiens 13-SEP-2001

Gaps ; 0 DB 1; Length 21; 1; Indels Score 16.4; DE Pred. No. 99; 0; Mismatches ò 3.8%; .48; Query Match
Best Local Similarity 94.4
Matches 17, Conservative

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ABZ58547 standard; DNA; 22 ABZ58547 RESULT S X S X Z X X Z X Z X X X A

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(first entry) 13-MAY-2003

ABZ58547;

primer X2R for detection of Fragile E site. PQ R Fragile E site; diagnosis; microcapillary electrophoresis; human; trinucleotide repeat; screening; PCR; primer; 88.

Homo sapiens.

WO2003014396-A1

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The present invention relates to a method for diagnosis of a multiplication disease of repeated trimucleotide sequence. The methods involves amplification of the repeated trimucleotide sequence by PCR, analysis of the amplified product on microcapillary electrophoresis (CE), analysis of the amplified product. In Fragile E site (FRAKE), in the size of the amplified product. In Fragile E site (FRAKE), in the all the size of the amplified product. In Fragile E site (FRAKE), in the healthy subjects and over 200 times in affected individuals. The present sequence is that of reverse primer X2R which is specific to the FRAKE repeated trimuclactide sequence region. It is used with forward primer X2R (see ABZ58546) to detect FRAKE. A diagnosis kit comprising these primers is claimed. In a healthy subject, a PCR product of 151 bp is produced. Use of CE, especially fabricated as an on-chip analysis system, allows the size of the PCR product to be measured rapidly, with accuracy and reproducibility. The method allows diagnosis before the disease cornot. It can be applied as a general screening test
                                                                                                                                                                                                                                                                       Diagnosing multiplication disease of repeated trinucleotide sequences e.g. Huntington's disease, by amplifying repeated trinucleotide sequence region, migrating and separating product by microcapillary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22 BP; 4 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                       Han S;
                                                                                                                                                                                                                                                                                                                                                                                     Claim 12; Page 8; 45pp; English.
                                                                                                                                                                                         Kim H,
                                                             06-AUG-2002; 2002WO-KR001489
                                                                                                  06-AUG-2001; 2001KR-00047301
                                                                                                                                                                                         Baik S,
                                                                                                                                                (BIOM-) BIOMEDLAB CORP
                                                                                                                                                                                                                                WPI; 2003-256603/25
                                                                                                                                                                                                                                                                                                                                              electrophoresis.
                                                                                                                                                                                         Lee Y,
                    20-FEB-2003.
                                                                                                                                                                                       Kim J,
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Gaps ö Score 16.2; DB 1; Length 22; Pred. No. 1.2e+02; 0; Mismatches 3; Indels 3.8%; Similarity 85.7 18; Conservative Query Match Local Matches

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86 Ŋ 66 CTGCACTACGAGGGCCGCGCA 22 ctrcceraceaeeeccececa

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BP. ADC58126 standard; cDNA; 24 (first entry) 18-DEC-2003 ADC58126; RESULT 48 ADC58126/c 

cancer; PCR; Mastocyte-specific guanine trinucleotidase 17.49 primer mastocyte-specific guanine trinucleotidase; 17.49; HIV; 88 primer;

Unidentified

CN1381569-A.

27-NOV-2002.

18-APR-2001; 2001CN-00112607

18-APR-2001; 2001CN-00112607.

(BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.

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Page

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Gaps

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Mao Y,
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The present invention provides sequences of antisense oligonucleotides targeted at the murine and human B7-1 and B7-2 coding and mRNA sequences. The antisense sequences have phosphorothicate backbones and some nucleotides are 2'-methoxyethoxy residues. The sequences can be used in the treatment of inflammatory and autoimmune disorders, including asthma, inventle diabetes mellitus, myasthenia gravis, Graves' disease, rheumatoid arthritis, allograft rejection, inflammatory bowel disease, multiple sclerosis, psoriaals, systemic lupus erythematosus, contact dermatitis, rhinitis, allergies and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel compound for diagnosing, preventing and treating immune disorders, comprising an oligonucleotide that specifically hybridizes with a nucleic acid sequence encoding B7 protein.
(systemic) lupus erythematosus, multiple sclerosis, contact dermatitis, rhinitis, allergy, cancer and metastases. The oligonucleotides may also be used to manipulate T cell activation ex vivo; to determine or detect B7 protein expression; for diagnosis; as assay and purification reagents and to study physiological roles of B7 proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, mouse; B7-1; B7-2; antisense; PCR primer; inflammation; autoimmune disorder; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
3.7%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 1.2e+02;
Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                        3.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.2e+02; cive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human B7-1 mRNA antisense oligonucleotide SEQ ID NO: 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                          Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Karras JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sxample 1; Page 45; 162pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               398 GAAGGICTICTACGIGAIC 416
                                                                                                                                                                                                                    398 GAAGGICTICTACGIGAIC 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 GAGGGTCTTCTACGTGAGC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00326186
                                                                                                                                                                                                                                                                                                                             :829/c
AAF32829 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vickers TA,
                                                                                                                                                               Local Similarity 89.5
Les 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-049991/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                 23-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                             AAF32829;
                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                          RESULT 50
AAF32829/c
                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; human; B7; T cell; inflammation; autoimmune disease; cell activation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The oligonucleotides which specifically hybridise to B7 modulate its expression (and thus T cell activation and proliferation). This is particularly useful for treatment and prevention of inflammation and autoimmune diseases, e.g. asthma, (juvenile) diabetes, myasthenia gravis, Grave's disease, rheumatoid arthritis, allograft rejection, psoriasis,
                                                                                                                                                                                The invention relates to a novel mastocyte-specific guanine trinucleotidase 17.49. The protein is useful for treating diseases such as cancer and HIV infection. The current sequence represents a primer related to the mastocyte-specific guanine trinucleotidase 17.49 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligo:nucleotide(s) that modulate expression of B7 proteins - used for, e.g. controlling activation and proliferation of T cells, particularly for treatment, diagnosis and prevention of inflammation.
                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                          Polypeptide-mastocyte-specific guanine trinucleotidase-17.49 and polynucleotide for coding it.
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                                                                                                                                                                                                                                                                                                                           Length 24;
                                                                                                                                                                                                                                                                                                                       3.8%; Score 16; DB 1; Length 24;
79.2%; Pred. No. 1.6e+02;
ive 0; Mismatches 5; Indels

    .20
    /*tag= a
    /note= "Phosphorothioate linkages"

                                                                                                                                                                                                                                                                                           Sequence 24 BP; 3 A; 7 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human B7-1 targetted oligonucleotide 13801.
                                                                                                                                                Example 3; SEQ ID NO 3; 32pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                  GAGGGCGCGCAGTGGACATCACC 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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AAV47987 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                            Local Similarity 79.2
Les 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-387783/33.
                                                            WPI; 2003-249033/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell proliferation.
                                                                                                                                                                                                                                                         of the invention.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-DEC-1997;
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Matches
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Human; calreticulin; antisense compound; hyperproliferative disorder; cancer; autoimmune disease; viral infection; cardiovascular disease; antisense therapy; cytostatic; immunosuppressive; virucide; antisense; phosphorothioate backbone; ss.
                                                        Human calreticulin antisense oligonucleotide, ISIS 109305.
                                                                                                                                                                                                                                                                     *tag= c
mod_base= OTHER
note= "2'methoxyethyl nucleotides"
                                                                                                                                                                                              "mod_base= OTHER
note= "2'methoxyethyl nucleotides"
                                                                                                                                                              mod_base= OTHER
note= "Phosphorothicate backbone"
                                                                                                                                      ocation/Qualifiers
                                                                                                                                                                                                                                             /*tag= e
/mod_base= m5c
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/mod_base= m5c
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mod_base= m5c
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mod_base= m5c
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mod_base= m5c
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mod_base= m5c
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mod_base= m5c
AAD39512/c
ID AAD39512 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-OCT-2001; 2001WO-US049045.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-OCT-2000; 2000US-00702327
                                                                                                                                                                                                                                                                                                                                                                            /*tag= i
/mod_base= n
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*tag= a
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                                        04-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-479759/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200236743-A2
                                                                                                                                      Key
modified_base
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                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-MAY-2002
                                                                                                                        Synthetic
                        AAD39512;
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The invention relates to antisense compounds, compositions and methods antisense computations the expression of calreticulin. The compositions comprise antisense compounds, particularly antisense oligonucleotides, targetted to nucleic acids encoding calreticulin. The antisense compound is useful for inhibiting the expression of calreticulin in human calls or tissues. It is also useful for treating a human having a disease or condition associated with calreticulin, e.g., hyperproliferative disorder e.g. cancer, autoimmune disease, viral infection or cardiovascular disease, by inhibiting expression of calreticulin. It is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. It is also used in antisense therapy. The present sequence is an antisense compound targetted to human calreticulin. This sequence is used to study the antisense inhibition of calreticulin expression-phosphorothioate 2'-MOE gapmer oligonucleotides
                            Claim 3; Page 82; 109pp; English.
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Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;

Gaps . 0 3.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred, No. 1.2e+02; tive 0; Mismatches 2; Indels Local Similarity 89.5 les 17; Conservative Query Match Best Loc Matches

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CCGAGGGCTGGGACGAAGA 43 19 CCGAGGACTGGGATGAGA 1 25 g δ

Human oligonucleotide sequence. ABZ92967 standard; DNA; 20 (first entry) 17-0CT-2003 ABZ92967; RESULT 52 ABZ92967 ID ABZ9

Human, antisense; lung dysfunction, nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

23-APR-2002; 2002WO-US013135. WO200285308-A2. Homo sapiens. 31-OCT-2002. 

E, Pabalan J, Katz 1 S; Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, 1 Miller S,

24-APR-2001; 2001US-0286137P.

(EPIG-) RPIGENESIS PHARM INC

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 8209; 872pp; English.

which has a The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions,

891. BIII

Novel antisense compound targeted to nucleic acid encoding calreticulin, useful for treating a human having disease or condition associated with calreticulin e.g. cancer, viral infection, autoimmune disease.

Wed Apr 41 14:36:41 400%

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Wed Apr 41 12:58:41 2004
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junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therappy. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of cof, or reducing sensitivity to adenosine, reducing levels of denosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences 

Seguence 20 BP; 0 A; 8 C; 8 G; 4 T; 0 U; 0 Other;

Gaps . 0 3.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.2e+02; live 0; Mismatches 2; Indels Local Similarity 89.5 nes 17; Conservative Query Match fatches

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131 GCTGGCCCCGCCTGGCGGTG 149 derecededechedenere 20

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851/c ADC65851 standard; DNA; 20 RESULT 53 ADC65851/

ВP

18-DEC-2003 (first entry) ADC65851;

Mouse TGF-beta receptor II targeted antisense oligonucleotide #50.

mouse, antisense oligonucleotide, transforming growth factor beta receptor II; TGF-beta receptor II; typerproliferative disorder, breast cancer; autoimmune disorder; rheumatoid arthritis; 2.0-merhoxyethyl gapmer; phosphorothioate backbone; ss; murine.

Mus musculus.

WO2003000656-A2.

03-JAN-2003.

19-JUN-2002; 2002WO-US019665.

21-JUN-2001; 2001US-00888361.

(ISIS-) ISIS PHARM INC

Murray SF, Wyatt JR;

WPI; 2003-175279/17.

New compound having a sequence targeted to a nucleic acid encoding Transforming growth factor beta-receptor II, useful for preparing a composition for treating hyperproliferative disorder e.g., lung, liver, colon or gastric cancer.

Claim 3; SEQ ID NO 147; 141pp; English.

The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. rheumatoid arthritis). The present DNA sequence

represents a 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to mouse TGF-beta receptor II. Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other; 88866

ö Query Match
3.7%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 1.2e+02;
Matches 17; Conservative 0; Mismatches 2; Indels

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RESULT 54 ADE27764/

踞 7764/c ADE27764 standard; DNA; 20

ADE27764;

(first entry) 29-JAN-2004 Human B7-1 mRNA targeted oligonucleotide SEQ ID 26.

ss; human; B7-1; inflammatory skin disorder; antisense; psoriasis; contact dermatitis; atopic dermatitis; seborrheic dermatitis; nummular dermatitis; generalised exfoliative dermatitis; eczema; critical costimulatory molecule.

Hômo sapiens. Synthetic.

US2003176374-A1.

18-SEP-2003.

09-MAY-2001; 2001US-00851871

31-DEC-1996; 96US-00777266. 04-UUN-1999; 99US-00326186. 25-MAY-2000; 2000WO-US014471.

(VICK/) VICKERS T A. (KARR/) KARRAS J G.

Karras JG; Bennett CF, Vickers TA,

WPI; 2003-863863/80.

Treating an inflammatory skin disorder such as psoriasis comprises topically applying an antisense compound targeted to the nucleic acid encoding human B7 protein.

Example 1; SEQ ID NO 26; 88pp; English.

The invention relates to a method of treating an inflammatory skin disorder in an individual by topically applying an antisense compound targeted to a nucleic acid molecule encoding a human B7 protein. The invention is for treating an inflammatory skin disorder in individual. The skin disorder is psoriasis, contact dermatitis, atopic dermatitis, seborrheic dermatitis, nummular dermatitis, generalised exfoliative dermatitis or eczema. The invention effectively modulates critical costimulatory molecules such as the B7 protein. The present sequence represents a human B7-1 targeted oligonucleotide. 

Sequence 20 BP; 5 A; 8 C; 4 G; 3 T; 0 U; 0 Other;

ö Query Match 3.7%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 1.2e+02; Matches 17; Conservative 0; Mismatches 2; Indels

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Gaps

398 GAAGGTCTTCTACGTGATC 416

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autoimmune disorder; immune disorder;

MUC-1; immunosuppression; autoimmune di inflammatory disorder; PCR primer; ss.

WO200034468-A2. Homo sapiens

(first entry)

06-NOV-2000

AAA63180;

Human muc-1 PCR primer #2.

99WO-US029016 98US-0111973P

09-DEC-1999; 11-DEC-1998;

15-JUN-2000.

Agrawal B, Longenecker BM;

(BIOM-) BIOMIRA INC.

WPI; 2000-423418/36.

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AAA63180 standard; DNA; 21

**AAA**63180

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A method has been developed for detecting T-cell activation by evaluating the amount of MUC-1 mucin expression in a T-cell compared to a non-darded control. The method is useful for treating disorders associated with T-cell activation, using an agent intrody/antagonist) that modulates MUC-1 activity. The T-cell activation associated disorders may be utoimmune or inflammatory disorders (e.g. inflammatory arthritis, portasis, allergies, allergic contact dermatis, rhemmatoris and arthritis, psoriasis, allergies, allergic contact dermatis, ankylosing spondylitis, myasthenia gravis, systemic lupus erythematosus, poblyarteritis nodosa, Goodpastures syndrome, isopathic thrombocytopenic purpura, autoimmune haemolytic anaemia, Grave's disease, rhemmatic fever, permicious anaemia, insulin-resistant diabetes mellitus, bullous permicious anaemia, insulin-resistant diabetes mellitus, bullous pemphigus vulgaris, viral myocarditis (Cocksakie B virus response), autoimmune thyroiditis (Hashimoto's disease), male infertility (autoimmune), sarcoidosis, allergic encephalomyelitis multiple scherosis, Sjorgens disease, Reiter's disease, Celiac disease, cancer. The present sequence represente a PCR primer for human MUC-1, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting T-cell activation by measuring the amount of MUC-1 expression useful for diagnosing or treating autoimmune or inflammatory disorders, viral disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; MUC-1; detection; T-cell activation; mucin; antiinflammatory; immunomodulator; antixheumatic; antiathritic; antiallergic; dermatological; antidabetic; nephrotropic; antithyroid; antianaemic; neuroprotective; hepatotropic; uropathic; ophthalmological; antiviral; cytostatic; autoimmune disorder; inflammatory disorder; viral disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 4 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 21; 40pp; English.
                                                                                                                                                                                                                                                                    BP
19 GAGGGTCTTCTACGTGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99WO-US012820.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Agrawal B, Longenecker BM;
                                                                                                                                                                                                                                                                    AAZ91293 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human MUC-1 PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BIOM-) BIOMIRA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-170935/15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                  AAZ91293;
                                                                                                                                                                                  RESULT 55
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Use of agent capable of intracellularly inhibiting mucin MUC-1 for inducing T-cell-based immunosuppression and for treating autoimmune disorders, transplant rejection and inflammatory disorders.

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The present sequence is a PCR primer for the human muc-1 mRNA. It was used to amplify the sequence in order to determine the expression pattern of the protein. This showed that MUC-1 is a fimunosuppressor, and its antagonists act to reduce overactive immune responses. Thus, MUC-1 required to treat inflammatory disorders such as rheumaticia arthritis, psoriasis, allergic contect dermatitis and ankylosing spondylitis, autoimmune disorders including myasthenia gravis, systemic lupus erythematosus, polyarteritis nodosa, Goodpastures syndrome, isopathic thrombocytopoenic purpura, autoimmune haemolytic anaemia, Graves' disease, rheumatic fever, pernicious anaemia, insulinersistant myocarditis, autoimmune thyroiditis, male infertility, sarcoidosis, viral myocarditis, autoimmune thyroiditis, male infertility, sarcoidosis, allergic encephalomyelitis, multiple sclerosis, Sjorgens disease, and primary cirrhosis, immune disorders, graft versus host disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.7%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 1.3e+02; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human IGERB coding sequence PCR primer SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 4 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 35; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       231 AAATCGGGAGGCTGCTTCC 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATATCGAGAGGCTGCTTCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              멾
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF92239 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transplant rejection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-MAY-2001
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Query Match 3.7%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 1.3e+02; Matches 17; Conservative 0; Mismatches 2; Indels

231 AAATCGGGAGGCTGCTTCC 249 ATATCGAGAGGCTGCTTCC 21

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                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein and coding sequences of several polymorphic variants of the human immunoglobulin E receptor beta chain (IGERB). These contain single nucleotide polymorphisms (SNPB) which may be indicative of a predisposition to atopy, allergy, asthma, rhinitis and eczema. Also provided are the sequences of probes and primers for use in identifying the genotype of an individual with regards to the IGERB gene. The IGERB gene is found at human chromosome 11q13. The sequences are all useful in therapeutics. The present sequence was used to isolate
Human; immunoglobulin E receptor beta chain; IGERB; chromosome 11q13; allergy; asthma; rhinitis; eczema; single nucleotide polymorphism; SNP; atopy; probe; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene construct; genome modification; higher plant; plant; marker gene; homologous recombination; cloning site; T-DNA; plant transformation; monocotyledon; Agrobacterium; gene function analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                        Novel polynucleotide useful for therapeutic purposes, comprises nucleotide polymorphisms in immunoglobulin E receptor beta chain gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.7%; Score 15.6; DB 1; Length 22; 81.8%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 2 A; 7 C; 5 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                               Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31 GCTGGGACGAAGATGGCCACCA 52
                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 77; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                  GENA-) GENAISSANCE PHARM INC
                                                                                                                                                       11-AUG-2000; 2000WO-US022175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-AUG-2002; 2002WO-JP008506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8-AUG-2001; 2001JP-00258489
                                                                                                                                                                                     99US-0150423P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACF03722 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NISB ) JAPAN TOBACCO INC. (SYGN ) SYNGENTA LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18; Conservative
                                                                                                                                                                                                                                                               Denton RR, Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer WxR-R3831.
                                                                                                                                                                                                                                 NAND/) NANDABALAN K.
                                                                                                                                                                                                                                                                                            WPI; 2001-226623/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VO2003020940-A1
                                                                                             WO200114588-A1
                                                                                                                                                                                    24-AUG-1999;
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                                                                                                                           01-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACF03722;
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The present invention describes a gene construct (1) for modifying the genome of higher plants by homologous recombination. (1) comprises marker genes and cloning sites between the right bottom sequence (BR) and left bottom sequence (BR) and left in the sequence (BR) and left constructs, particularly with a first cloning site for integration into the 5' region in the monologous recombination of the target gene into the 5' region in the homologous recombination of the target gene into the 3' region in the homologous recombination of the target gene into the 3' region in the homologous recombination of the target gene into the 3' region in the homologous recombination of the target gene into the nest genome, and a second cloning site for integration into the bost genome, and (2) producing a genome-modified higher plant (especially a monocotyledon) by using homologous recombination comprising (1) infecting plant cells, tissues or calluess with the Agrobacterium, (11) infecting plant cells, tissues or calluess produced by homologous recombination through negative or positive selection; (iv) culturing selected cells or tissues into calluess; (v) culturing in callue. Regenerating medium to grow into heterozygously modified plants; and (vi) producing homozygously modified plants; and (vi) producing homozygously modified plants; are useful for modifying the genome of higher plants by monologous recombination without altering the original locus, for the analysis of genomical dynamics. The present sequence represents a PCR primer which is used in an example from the present invention
                                                                                                      A gene construct for modifying the genome of higher plants by homologous recombination without altering the original locus, comprises marker genes and cloning sites between the right and left bottom sequences from T-DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; anti-ckel; antisense oligonucleotide; benign lesion; papilloma; atherosclerosis; psoriasis; autoimmune disease; bacterial infection; viral infection; HIV; hepatitis; herpes; polythemia; mastocytosis; cksl inhibitor; skp2 inhibitor; cytostatic; antisense therapy; sarcoma; leukaemia; Hodgkin's lymphoma, non-Hodgkin's lymphoma; adenoma; melanoma; carcinoma; colon cancer; pancreatic cancer; cervical cancer; human; Skp2;
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22 BP; 2 A; 5 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide SEQ ID NO:40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          372 TTCCTGGACCGCGACGGCG 393
                                                                                                                                                                                                    Example 5; Page 22; 48pp; Japanese.
                     Inagaki Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 81.8%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA14342 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                       Terada R,
                                                              WPI; 2003-332936/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003068939-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-AUG-2003
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                       Iida S,
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12-FEB-2002; 2002US-0356906P.
         Reinhard C,
              WPI; 2003-689667/65.
    (CHIR ) CHIRON CORP
         Walter AO,
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The present invention describes a method for treating cancer comprising using an anti-cks1 antisense oligomucleotide. Also described: (1) treating benign lesions (e.g. papillomas, atherosclerosis and psoriasis), autoimmune diseases, bacterial infections, viral infections (e.g. HIV infections, hepatitis or herpes infections), polythemia, or mastocytosis using the cks1 antisense oligomucleotide or as skp2 inhibitor; (2) treating cancer using a skp2 inhibitor; (3) cks1 inhibitor; (4) a ribozyme, a protein, a polypeptide, an antibody or a small molecule; (4) in solated polynucleotide with a sequence comprising a transcriptional initiation region and a sequence encoding an antisense oligomucleotide; (5) a recombinant vector comprising the polynucleotide; and (6) inhibiting the expression of cks1 or skp2 in a mammalian cell. cks1 and csn be used in antisense therapy, and as Cks1 and Skp2 inhibitors. The method is used in antisense therapy, and as Cks1 and Skp2 inhibitors. The method is used in antisense melanomas, carcinomas, colon cancer, pancreatic cancer, or cervical cancer. The present sequence represents an antisense oligomucleotide given in the Sequence Listing of the present invention. Treating cancer, e.g. sarcoma, leukemia, (non-)Hodgkin's lymphoma, adenomas, melanomas, carcinomas, colon cancer, pancreatic cancer or cervical cancer, by employing an anti-cks-1 antisense oligonucleotide. Sequence 23 BP; 8 A; 9 C; 5 G; 1 T; 0 U; 0 Other; Disclosure; Page 86; 87pp; English.

3:7%; Score 15.6; DB 1; Length 23; 81.8%; Pred. No. 1.8e+02; tive 0; Mismatches 4; Indels GCGCAGTGGACATCACCACGTC 103 GCGCAGCAGACAAACCACGTC 22 18; Conservative Local Similarity 82 Query Match Best Loca Matches à 셤

Gapa . 0

> ABT03847 standard; DNA; 24 (first entry) 13-SEP-2002 ABT03847; RESULT 60 ABT03847

BP

Human RFC40kD gene PCR primer SEQ ID NO: 368

Human; cancer; neoplastic disease; tumour specific marker; cytostatic; transcription factor; PCR; primer; ss.

WO200240716-A2 Homo sapiens.

23-MAY-2002.

13-NOV-2001; 2001WO-US043461

16-NOV-2000; 2000US-0249508P

(CEMI-) CEMINES LLC

Palm K;

WPI; 2002-537346/57

Determining the presence of neoplastic molecular markers, by identifying the presence of markers in host test sample using array of neoplastic molecular marker specific reagents and analyzing the array of the reagents

21; 41pp; English. Page ; Example 1;

Shamoon BF

Jefferson AB,

븅 The present invention relates to a method for determining the presence oneoplastic molecular markers in a host, involving the use of neoplastic molecular marker specific reagents to detect such markers and analysing the array of reagents, allowing the identification of the meoplastic disease present. This can be used to determine the best treatment for cancers, in particular neural cell, lung and prostate tumours. The present sequence is a PCR primer useful for detecting the coding sequences of markers of the invention ####X#X00000000X8

Sequence 24 BP; 4 A; 6 C; 6 G; 8 T; 0 U; 0 Other;

Gaps .; 0 Length 24; 4; Indels 3.7%; Score 15.6; DB 1; 31.8%; Pred. No. 1.9e+02; Ive 0; Mismatches 4; Query Match 3.7%; Best Local Similarity 81.8%; Matches 18; Conservative

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72 24 51 CACTCAGAGGAGTCTCTGCACT cagicagraaagicicigcici

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ABL54647 standard; DNA; 17 ABL54647; ABL54647 

BP.

(first entry) 31-MAY-2002 Human p53AIPI associated PCR primer SEQ ID NO 20.

Human, p53, p53AIPI, p53-dependent apoptosis-associated, apoptosis; cytostatic, cancer; PCR, primer, ss.

Homo sapiens.

WO200212496-A1 14-FEB-2002. 02-AUG-2001; 2001WO-JP006666.

03-AUG-2000; 2000JP-00240399

(UYTY ) UNIV TOKYO. (ONCO-) ONCOTHERAPY SCI INC.

Nakamura Y, Arakawa H;

WPI; 2002-217192/27.

p53-dependent apoptosis-associated protein and its encoding gene p53AIPI, used for screening apoptosis mediated remedies for cancer and as controllers of apoptosis induction.

Example 7; Page 40; 121pp; Japanese.

The invention relates to human p53-dependent apoptosis-associated protein, P53AIPI comprising fully defined 806, 777, 2659 nucleotide sequences (ABL54631-ABL54633 respectively) given in the specification and proteins having fully defined 124, 86 and 108 amino acid sequences (ABB08837-ABB08839 respectively) given in the specification. The protein and encoded gene have cytostatic activity, are useful in screening for regulators of apoptosis for subsequent use as cancer treatments. The present sequence is that of the Human p53AIPI associated PCR primer, useful to the invention

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RESULT 62 AAX38484/

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The present invention relates to an expression library comprising synthetic nucleic acid sequences, not cloned from an immunized source, where the nucleic acid sequences are derived from an mutagenised immunoglobulins that are naturally devoid of light chains. The library is useful for the preparation of antibodies having binding specificity for a target antigen which avoids the need for a donor to have been previously immunized with the target antigen. The recombination of heavy and light chains is avoided, therefore preventing the formation of molecules that are non-functional. The number of hypervariable residues in the binding domain is reduced, allowing a more complete repertoire of possible binding variants to be obtained. The present sequence is a PCF primer targeted to anchor regions in Ilama antibodies. The primers (AAA73745 to AAA73745 amplified the framework regions H, P2, P2, P2, P3 and P4. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Expression library comprising nucleic acids not cloned from an immunized source, derived from immunoglobulins naturally devoid of light chains, use for producing antibodies specific for a target antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Heavy chain; tetanus; toxin; human; monoclonal; antibody; K4.1; hybridoma; immortalisation; in vivo; xenomice; analysis; immunoglobulin; diagnosis; research; therapy; B cell; primer; polymerase chain reaction; amplification; PCR; ss.
                                                                                          Llama; primer; expression library; antibody; immunization; anchor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-tetanus toxin human antibody heavy chain cDNA primer MG-24Vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1; Length 20; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                   Primer F3 used to amplify part of llama antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 2 A; 7 C; 7 G; 3 T; 0 U; 1 Other;
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(UNIL ) UNILEVER NV.
(HIND-) HINDUSTAN LEVER LTD.
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(revised)
(first entry)
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Best Local Similarity 85.0
Matches 17; Conservative
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                                                                                                                framework; ss
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15-SEP-2003
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                                                                                                                                                     Lama glama.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               X38552) which are nuclease resistant, and comprises about 3-50 nucleotides complementary to the ribonucleotide reductase gene or the secA gene of a microorganism. The antisense oligonucleotides are used to treat mammalian pathological conditions mediated by microorganisms. The oligonucleotides are particularly useful as antimicrobial agents in crop
                                                                                                                                                                                                                                                                                                                                                                                                                     Microorganism inhibitor; antisense; nuclease resistant; treatment; ribonucleotide reductase; secA gene; pathological condition; R1 subunit; antimicrobial agent; crop protection; primer; R2 subunit; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S
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                                                       DB 1; Length 17;
                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 8 C; 3 G; 6 T; 0 U; 0 Other;
                    Sequence 17 BP; 2 A; 6 C; 2 G; 7 T; 0 U; 0 Other;
                                                       Score 15.4; DB
Pred. No. 98;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    E. coli SecA antisense oligonucleotide 40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 24; 103pp; English
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                                                                                                                                   206 GAAAGCAGAGAACTCGG 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA73747 standard; DNA; 20 BP
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                                                       3.6%;
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                                                                                                                                                                   GAAAGCAGAGAACTTGG
                                                                                                                                                                                                                                                                      AAX38484 standard; DNA; 20
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                                                       Query Match
Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-120874/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JUL-1998;
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                                                                                                                                                                                                                                                                                                           AAX38484;
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Best Local S
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Best Loca Matches

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RESULT 63
AAA73747/
ID AAA7
XX
AC AAA7

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The invention relates to modification of the half-life of an antibody.

The method of the invention comprises physically linking an antibody which contains an FCRn receptor binding molety, (hinge, CH2 and CH3 contains an FCRn receptor binding molety, (hinge, CH2 and CH3 molecules are protected from degradation by the endosomal FCRb/FCRn receptors, which gives them a relatively extended serum half-life or an antibody by increasing molety further extends the serum half-life of an antibody By increasing molety further extends the serum half-life of an antibody by increasing che serum half life of an antibody by increasing clinical treatments is lowered. This could significantly lower costs for treatment, and lead to less frequent hospital visits as fewer doses are required, thereby increasing the quality of life for patients, and potentially reducing the likelihood of toxicity. The technology can also be adapted to extend the serum half life of other proteins, in addition to antibodies. Sequences AAA66862-A66863 represent PCR primers used in an exemplification of the present invention to amplify cDNA generated from human monocolonal antibody poly(A+) mRNA expressed in XenoMice. The PCR products were then cloned into pCRII and sequenced
                                                                                  Modifying antibody half life by linking the antibody to an FcRn binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytotoxic T-lymphocyte antigen-4; CTLA-4; antibody; immune system; hyperimmunity disorder; autoimmune disease; diabetes; graft rejection; proliferative disorder; cancer; immunodeficient disorder; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 3.6%; Score 15.2; DB 1; Length 23; Best Local Similarity 81.0%; Pred. No. 2.1e+02; Matches 17; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         263 GGTGCACCTGGAGCAGGCGG 283
                                                                                                                                                    Example 1; Page 47; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 GGTGCAGCTGGAGCAGTCNGG 23
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    Foord O;
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    Junghans R,
                                             WPI; 2000-224282/19.
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modified_base
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    Gallo M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Antibody contg. fully human variable region specifically reactive with antigen - prepd. by immunisation of non-human animal incapable of \underline{p}roducing endogenous immunoglobulin (Ig), but capable of producing human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a primer for the PCR amplification of the CDNA encoding the heavy chain of the anti-teranus toxin (TT) human monoclonal antibody (MAD) K4.1, which was secreted by the hybridoma K4.1 and obtained by immortalising B cells from xenomice (containing integrated human DNA from the immunoglobulin locus) immunised with TT. The MAD can be used for analysis, diagnosis, research and therapy, particularly for human therapeutic, and in vivo diagnostic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IgG1; immunoglobulin G; FCRn receptor; FCRb; VH region; heavy chain variable region; serum half life; monoclonal antibody; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                  Brenner DG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                  Klapholz S,
                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             263 GGTGCACCTGGAGCAGGGCGG 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7; Page 28; 64pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA06862 standard; DNA; 23 BP.
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                                                                                                                                                                                                                                                                                                                      (CELL-) CELL GENESYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-497628/49.
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                                         Key
modified_base
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                                                                                                                                                                                                                                                                             28-APR-1995;
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Synthetic.
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Davis CG;

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Corvalan JR;

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Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                  Example 2; Page 66; 157pp; English
                                                                                                                            263 GGTGCACCTGGAGCAGGGCGG 283
                                                                                                                                                            ACD10944 standard; DNA; 23
                                                                                                        Query Match
Best Local Similarity 81.01
                                                                                                                                                                                                                                                                                                      WPI; 2003-328430/31
        WPI; 2000-442647/38.
                                                                                                                                                                                                                                 US2002173629-A1.
                                                                                                                                                                                                                                                                                              Takobovits A,
                                                                                                                                                                                                                                                  05-NOV-1998;
                                                                                                                                                                                                                                                           05-MAY-1997;
                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                29-SEP-1998;
                                                                                                                                                                             12-AUG-2003
                                                                                                                                                                                                                                          21-NOV-2002
                                                                                                                                                                     ACD10944;
                                                                                                                                                                                                                                                                                 (GALL/)
(JIAX/)
                                                                                                                                                                                                                                                                        JAKO/)
                                                                                                                                                                                                                                                                             YANG/)
                                                                                                                                                        ACD10944
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(first entry)

23

GCTGCAGCTGGAGCAGTCNGG

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Jia X;

Gallo M,

Yang X,

JAKOBOVITS A.

YANG X. GALLO M.

JIA X.

98US-00187693 97US-00851362. 98US-00162280. receptors, useful in cancer therapy.

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The invention relates to a novel chimeric or human monoclonal antibody or its antigen-binding portion that specifically binds to and activates human CD40. The anti-CD40 antibody of the invention demonstrates cytoatatic, virucide, antibacterial, immunossimulant and anti-HIV activities and may be useful for treating a hyperproliferative disorder such as cancer, viral and bacterial infection or genetic, primary or combined immunodeficiency conditions including neutropenia or HIV infection. The anti-CD40 antibodies may also be useful for detecting CD40 in a biological sample in vitro or in vivo, as well as during gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New chimeric or human monoclonal antibody or its antigen-binding portion that specifically binds to and activates human CD40, useful for enhancing an immune response in a human, or treating cancer, HIV, neutropenia or
                                                      The invention relates to an antibody that binds to an epidermal growth factor receptor (FGF-r) and exhibits inhibition of Expropriation of Exp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  anti-CD40 monoclonal antibody; CD40; cytostatic; virucide; antibacterial; immunostimulant; anti-HIV; hyperproliferative; cancer; viral; bacterial infection; immunodeficiency; neutropenia; HIV; gene therapy; human; PCR; primer; ss; universal; VH; MG-30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.6%; Score 15.2; DB 1; Length 23; 81.0%; Pred. No. 2.1e+02; tive 0; Mismatches 4; Indel8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jia X,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Universal human VH PCR primer MG-30.
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Example 3; Page 17; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   263 GGTGCACCTGGAGCAGGGCGG 283
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 81.0
es 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE28495 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bedian V, Gladue RP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-441521/41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2004
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                                                                                                                                                                                                                                                                                                                                  The present sequence represents a PCR primer which is used to amplify a fragment of the gene encoding a heavy chain of human antibodies against cytotoxic T-lymphocyte antigen (CTAA)-4. The specification describes an synthetic antibody which is capable of binding CTLA-4. The antibody is composed of a heavy chain variable region, comprising a modified contiguous sequence from a FRI-FRS sequence encoded by a human VH3-33 family gene. The modifications are contained in CDR1, CDR2 and/or framework regions. The antibodies may be used to inhibit CTLA-4 and down-regulate the immune system to treat hyperimmunity disorders (e.g. autoimmune disease, diabetes and graft rejection) and proliferative disorders (e.g. cancer). CTLA-4 stimulatory agents may be used to upregulate immune system to up-regulate immunodeficient disorders
                                                                                                                                  Novel antibodies capable of binding cytotoxic T-lymphocyte antigen (CTLA) -4 containing specified heavy and light chain sequences, useful for treating, e.g. immune disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human epidermal growth factor receptor (EGF-r) antibody PCR primer #1.
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AAT32459;

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missing 5' sequence of cDNA encoding a novel canalicular multispecific organic anion transporter (cMOAT) protein, isolated from a human lambda gtil liver cDNA library. The protein is a new member of the ATP-binding cassette (ABC) transporter family. The ATP dependent cMOAT transporter system mediates hepatobiliary excretion in the liver. cMOAT may be a liver-specific homologue of multidrug resistance-associated protein. The nucleic acids are used to provide cells with cMOAT protein activity. CMOAT protein activity in cells can be enhanced by increasing the level of glutathions, glucuronide and/or sulphate. Antiesness constructs, especially derived from another multidrug resistance (WDR)-related protein, e.g. MDR-1, to the nucleic acids and vectors can be used to decrease the level of cMOAT in a cell. The nucleic acids and proteins can be used especially in diagnosis of Dubin-Johnson disease, Rotor disease or another disease involving cMOAT. The cMOAT gene may also be used as a selectable marker gene. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                        Canalicular multispecific organic anion transporter protein; for ATP-binding casette transporter family; ABC transporter; hepatobiliary excretion; multidrug resistance-associated protein; cMOAT protein activity; multidrug resistance-related protein; MDR-1; Dubin-Johnson disease; Rotor disease; PCR primer; ss.
                                                                                                                        used to isolate the missing 5' sequence of rat.cMOAT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA encoding human and rat canalicular multispecific organic transporter proteins - useful for diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15; DB 1; Length 23;
Pred. No. 2.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bosma PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      393 GCCAAGAAGGTCTTCTACGTGAT 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CENT AMSTERDAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 16; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HEIN-) HET NEDERLANDS KANKER INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Johnson disease and Rotor disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Paulusma CC,
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                                                                                 (first entry)
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                                                              (revised)
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                                                                                                                      PCR primer 2
                                                                                                                                                                                                                                                                                                                                                                                                                            21-FEB-1997;
                                                                                                                                                                                                                                                                                                                                             WO9731111-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-FEB-1996;
                                                            25-MAR-2003
                                                                                 01-APR-1998
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                                                                                                                                                                                                                                                                                                          Rattus sp.
                                                                                                                                                                                                                                                                                     Synthetic
                       AAT94027;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The calpain large subunit 1 gene located on chromosome 15 codes for a calcium activated neutral protease (CANP3) belonging to the calpain family. Mutations in the gene induce limb-girdle muscular dystrophy (LGMD) 2 disease. The gene, and fragments of it, can be used in the prevention, treatment, diagnosis and detection of a predisposition to CAMP2 disease. Bight primers (AAT32456-63) were used to localise the calpain large subunit 1 gene. The results positioned the gene in a region previously defined as 15q15.1-q21.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human novel Calpain large sub:unit 1 gene encoding a calcium dependent protease - used to develop prods. for the diagnosis and treatment of limb -girdle muscular dystrophy 2 disease.
therapy procedures. The current sequence is that of the human anti-CD40 antibody-related PCR primer of the invention.
                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Calpain, subunit, calcium, protease, mutation, treatment, detection, identification, diagnosis, limg girdle muscular dystrophy, LGMD2, calcium activated neutral protease, CANP, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer (P94in13) for localisation of calpain large subunit 1 gene.
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                                                                                                  Length 23;
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Pred. No. 1.9e+02;
Mismatches 0; Indels
                                                                                                                                           4; Indels
                                                            Sequence 23 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 1 Other;
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                                                                                                Score 15.2; DB 1;
Pred. No. 2.1e+02;
); Mismatches 4;
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                                                                                                                                                                                263 GGTGCACCTGGAGCAGGCGG 283
                                                                                                                                                                                                          3.5%; Sco.
100.0%; Pre
0;
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                                                                                                3.6%;
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                                                                                            Query Match
Best Local Similarity 81.0
Matches 17; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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AAD25481

BP.

AAT94027 standard; cDNA; 23

RESULT 70 AAT94027/C ID AAT94

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(ORTH ) ORTHO-MCNEIL PHARM INC.
                    Burris TP,
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Rybczynski PJ;

Benzoxazinone derivative, glucose metabolism, lipid metabolism, NIDDM; PPAR gamma; peroxisome proliferator activated receptor gamma; therapy; non-insulin dependant diabetes mellitus; nephropathy; neuropathy; FCOS; atherosclerosis; retinopathy; polycystic ovary syndrome; hypertension; ischaemia; obesity; heart disease; irritable bowel disorder; stroke; reduced insulin sensitivity; inflammation; cataract; aP2 mRNA; probe; ss. Probe #18 used in aP2 assay for antagonist. 12-MAY-2000; 2000US-0203859P. 11-MAY-2001; 2001US-00853798. 11-MAY-2001; 2001WO-US015320. 26-MAR-2002 (first entry) WO200187860-A2. Unidentified. 22-NOV-2001,

Use of benzoxazinone derivatives for treating a subject suffering from a disorder in glucose and lipid metabolism such as non-insulin dependant diabetes mellitus or obesity. WPI; 2002-082970/11.

Example 2; Page 34; 45pp; English.

The invention relates to benzoxazinone derivatives useful as peroxisome proliferator activated receptor (PPAR) gamma modulators. The invention daso relates to pharmaceutical compositions comprising benzoxazinone derivatives and methods for treating the onset of a disorder in glucose and lipid metabolism, preferably a condition of reduced insulin sensitivity such as non-insulin dependant diabetes mellitus (NIDDM), polycystic ovary syndrome (POCS), hypertension, ischaemia, stroke, heart diseases, irritable bowel disorder, inflammation and cataract. The present sequence is a probe designed to anneal to the aPZ mRNA and function in the bDNA mRNA decetion system. This probe used in the aPZ assay for antagonist which is used in the exemplification of the

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Sequence 23 BP; 5 A; 6 C; 4 G; 8 T; 0 U; 0 Other;

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Gaps
Match 3.5%; Score 15; DB 1; Length 23; Local Similarity 78.3%; Pred. No. 2.3e+02; les 18; Conservative 0; Mismatches 5; Indels
        Query Match
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TCTACAGCGACTTCCTCACTTTC 374 rciecaereacricercaaric 23 352

8 용

AAI68021 standard; DNA; 23 AAI65021; AAI6802 RESULT 

ВР

4h-Benzo(1,4)oxazin-3-one compound; glucose metabolism; lipid metabolism; peroxisome proliferator activated receptor; therapy, NIDDM; non-insulin dependant diabetes mellitus; nephropathy; neuropathy; NIDDM; atherosclerosis; retinopathy; polycystic ovary syndrome; hypertension; ischaemia; obesity; heart disease; irritable bowel disorder; cataract; anorectic; nephrotropic; ophthalmological; cytostatic; hypotensive; vasorropic; cerebroprotectiv; cardiant; antiinflammatory; probe;

Probe #18, used in aP2 assay for antagonist.

(first entry)

12-MAR-2002

AAD24705;

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AAD24705 standard; DNA; 23

RESULT 73 AAD24705

(first entry)

13-MAR-2002

ap2 mRNA specific oligonucleotide probe.

Peroxisome proliferator activated receptor gamma; benzoxazinone; NIDDM; non-insulin dependant diabetes mellitus; antidiabetic; anorectic; nephrotropic; ophthalmological; antiarteriosclerotic; cytostatic;

WO200187862-A2

aP2 mRNA; ss. Unidentified.

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The invention provides methods of treating a subject suffering from a condition associated with peroxisome proliferator activated receptor gamma (PpsAgamma) activity that involves administering a benzoxazinone compound of a specified formula to the subject. The method is useful for treating and inhibiting in a subject the onset of a condition associated with PpsAgamma activity such as a condition of reduced insulin sensitivity, non-insulin dependant diabetes mellitus, obesity, nephropathy, retinopathy, atherosclerosis, polycystic ovary girdrom, hypertension, ischemia, stroke, heart diseases, irritable bowel disorder, inflammation and cataract. Sequences AAI68004-023 represent oligonucleotide probes specific for ap2 used in ap2 mRNA assay for
                                                                                                                                                                                                                                                                                                                                                               Use of benzoxazinone derivatives for treating a subject suffering from a condition associated with peroxisome proliferator activated receptor gamma activity e.g. non-insulin dependant diabetes mellitus and obesity.
hypotensive; vasotropic; cerebroprotective; cardiant; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                             Turchi IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5; Indels
                                                                                                                                                                                                                                                                                             Combs DW, Rybczynski PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 5 A; 6 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.5%; Score 15; DB 1; I
ilarity 78.3%; Pred. No. 2.3e+02;
Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 7; Page 29; 46pp; English.
                                                                                                                                                                                                                                                         (ORIH ) ORTHO-MCNEIL PHARM INC
                                                                                                                                                                                                   12-MAY-2000; 2000US-0203861P.
11-MAY-2001; 2001US-00854368.
                                                                                                                                                                  11-MAY-2001; 2001WO-US015377
                     PPARgamma; probe; ap2; ss.
                                                                                                                                                                                                                                                                                               Ę
                                                                                                                                                                                                                                                                                             Burris TP, Demarest
                                                                                                                                                                                                                                                                                                                                WPI; 2002-082971/11.
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les 18; Conserv
                                                                                           WO200187861-A2
                                                                                                                              22-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antagonists
                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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22-NOV-2001

Yang XB;

Herzog N,

Luxon BA,

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monitoring biological interaction, modified aptamer; phosphorothioate agonist; phosphorothioate antagonist; antibacterial; phosphorothioate antagonist; antibacterial; antianthritic; antianthritic; antianthritic; corpusatio; anti-HIV; antianthritic; antianthritic; antianthritic; antianthritic; croin's disease; NF-kappaB; toxic shock; sepsis; rheumatory bowel disease; asbestos lung disease; Hodgkin's disease; prostate cancer; ventilator induced lung injury; cancer; AIDS; human cutanecus T cell lymphoma; lymphoid malignancy; HILV-1 induced adult T-cell leukaemia; atherosclerosis; cytomegalovirus; herpes simplex virus; UCV; 8V-40; rhinovirus; influenza;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide duplex Seq ID94 related to biological interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD43640 standard; DNA; 23 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003050290-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD43640;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD43640
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(first entry)

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This invention relates to a novel apparatus for monitoring biological interaction which comprises a substrate and a modified aptamer attached to the substrate, where a target molecule or its portion, contacted with the modified aptamer under conditions to allow formation of a complex between the modified aptamer and the target molecule or its portion, is detected. The invention may be useful in developing phosphorotrhioare agonists or antagonists which may have antibacterial, immunosuppressive, antitrheumatic, antiarthritic, antiinflammatory, cytostatic, anti-HIV, and apparatus of the present invention are useful for monitoring biological interactions and in functional proteomics. As an example, and apparatus of the present invention are useful for monitoring biological interactions and in functional proteomics. As an example, and apparatus of the present invention are useful for monitoring biological interactions and in functional proteomics. As an example, and apparatus of the present invention are useful for monitoring biological authritis, Crohn's disease, such as toxic shock, sepsis, returnating NF-kappaB aptamer-related diseases, such as toxic shock, sepsis, remained at third induced lung injury, general cancer, AIDS, human cutaneous T cell lymphoma, lymphoid mallghancies, HTM-1 induced adult T-cell leukaemia, atherosclerosis, cytomegalovirus, herpes simplex virus, JCV, SV-40, rhinovirus, influenza, neurological disorders and lymphomas. The present sequence is that of an oligomuclecide duplex which was used during the exemplification of the invention.
                                                                                                                                                                                                                                                                                              New apparatus with a substrate and a modified nucleotide aptamer for monitoring biological interactions, useful for diagnosing and treating NF-kB aptamer-related diseases, such as toxic shock, rheumatoid arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 23 BP; 1 A; 6 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 58; SEQ ID NO 94; 67pp; English.
                                 15-NOV-2001; 2001US-0334887P
                                                                                               (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                       WPI; 2003-513977/48.
                                                                                                                                                                                                                                                                                                                                                                                                             cancer and AIDS.
                                                                                                                                                                     Gorenstein D,
The patent discloses 4h-Benzo(1,4) oxazin-3-one compounds which are useful as peroxisome proliferator activated receptor (PPAR) gamma agonists and antagonists. The invention also relates to compositions comprising such compounds and methods for treating or inhibiting the onset of a disorder in glucose and lipid metabolism, preferably a condition of reduced in sulin sensitivity, such as non-insulin dependent diabetes mellitus (NIDDM), obesity, atherosclerosis, nephropathy, neuropathy, retinopathy, polycystic ovary syndrome, hypertension, ischaemia, stroke, heart diseases, irritable bowel disorder, inflammation and cataract. The present DNA sequence is a probe which is designed to anneal to ap2 mRNA and function in the bDNA mRNA detection system. This probe is used in ap2 assay for antagonist in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                          Use of 4h-benzo(1,4)oxazin-3-one derivatives for treating a subject suffering from a disorder in glucose and lipid metabolism e.g. non-insulin dependant diabetes mellitus and obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.5%; Score 15; DB 1; Length 23; Best Local Similarity 78.3%; Pred. No. 2.3e+02; Matches 18; Conservative 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 5 A; 6 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                         Burris TP, Combs DW, Rybczynski PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       352 TCTACAGCGACTTCCTCACTTTC 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 rergeadraderregreanire 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 38; Page 58; 76pp; English.
                                                                                                                                                                                                                                       (ORTH ) ORTHO-MCNEIL PHARM INC
                                                              11-MAY-2001; 2001WO-US015383.
                                                                                                                            12-MAY-2000; 2000US-0203860P.
11-MAY-2001; 2001US-00854302.
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                                                                                                                                                                                           G-alpha-13; human; inhibitor; cancer; antisense compound; therapy; ss.
                    Gaps
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3.5%; Score 15; DB 1; Length 23; 78.3%; Pred. No. 2.3e+02;
                   5; Indels
                                                                                                                                                                        Human G-alpha-13 antisense inhibitor ISIS# 20741.
                    0; Mismatches
                                         132 CTGGCCCGCCTGGCGGTGGAGGC 154
                                                           crerrccaecredcedrededec 23
                                                                                                              AAZ31792 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                       98US-00205860.
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                                                                                                                                                      (first entry)
Query Match
Best Local Similarity 78.3
Matches 18; Conservative
                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                      04-DEC-1998;
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                                                                                                                                                      24-JAN-2000
                                                                                                                                                                                                                                              US5981732-A.
                                                                                                                                                                                                                                                                   09-NOV-1999
                                                                                                                                                                                                                Synthetic
                                                                                                                                 AAZ31792;
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AAZ31792
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(ISIS-) ISIS PHARM INC

07-AUG-2002; 2002WO-US025049.

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E. coli ilvC gene PCR primer panE2.
                                                                                                                                                                                                                                                                                                                                        Example 4; Page 9; 24pp; German.
                                   Claim 11; Col 38; 38pp; English
                                                                                                                                                                                                                                                                                        Elischewski F, Kalinowski J,
Farwick M, Thierbach G;
                                                                                                                                 1 cacceccaceccaccaca 18
                                                                                                                                                           AAA12163 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                    98DE-01046499.
                                                                                                                      105 GACCGCGACCGCAGCAAG
                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                              DEGS ) DEGUSSA-HUELS AG
                                                                                                                                                                                                                                                                                                        WPI; 2000-304637/27.
              WPI; 1999-633376/54.
                                                                                                                                                                                                                          Escherichia coli.
                                                                                                                                                                                                                                    DE19846499-A1.
                                                                                                                                                                                                                                                          09-OCT-1998;
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                                                                                                                                                                                10-AUG-2000
                                                                                                                                                                                                                                               30-APR-2000
     Cowsert LM,
                                                                                                                                                                      AAA12163;
                                                                                                                                                 RESULT 76
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nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase (also known as PBPCK-mitochondrial phosphoenolpyruvate carboxykinase (also known as PBPCK-mitochondrial) PBPCK-M; PCKS and murpEPECK), where the Oligonucleotide specifically hybridise with and inhibit the expression of human mtPBPCK. The antisense oligonucleotides can be used for inhibiting the expression of mtPBPCK in human cells or particularly a human suspected of having or treating an animal, particularly a human suspected of having or being prone to a condition or disease associated with expression of mtPBPCK. They can also be used in diagnostics and as research reagents in sandwich and other assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound targeted to a nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase useful for treating a human with a mitochondrial phosphoenolpyruvate carboxykinase-associated
amplify the ilvC gene, panE gene, panB gene, panC gene, panD gene and the avtA gene which are used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, mitochondrial phosphoenolpyruvate carboxykinase; PEPCK-M; PCK2; PEPCK-mitochondrial; mtPEPCK; antisense oligonucleotide; modulation; phosphorothioate; inhibition; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mtPBPCK phosphorothioate antisense oligonucleotide SEQ ID NO:11.
                                                                                                                                                          Gaps
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                                                                                                            Length 20;
                                                                                                                                                        2; Indels

    .20
    /*tag= a
    /note= "phosphorothioate linkages"

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                                                                  Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                            Score 14.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Col 39; 32pp; English.
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                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-00366257.
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                                                                                                                   3.5%;
                                                                                                                                                                                                                                               3 Agrererreactaceage
                                                                                                                                                                                                    61 AGTCTCTGCACTACGAGG
                                                                                                                                                                                                                                                                                                                                                          AAZ95323 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                 3.5%
Query Match
Best Local Similarity 88.9%
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-205209/18.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                                   AAZ95323;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                       AAZ95323
                                                                                                                                                                                                                                                                                                                  RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel method for the production, and improvement, of panthothenic acid (I)-producing microorganisms by amplifying (particularly overexpressing) sequences (I) that encode ketopanthoate reductase (XRP), specifically the pans gene, either individually or together. Optionally the ilvC gene is also amplified. (I) is a vitamin used in cosmetics, medicine and human or animal nutrition. The method provides increased yields of (I), e.g. 35-40 mug/ml for the most productive strains. AAA12160-A12171 represent PCR primers used to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         useful as
that
                                                                                                                                                                                                    This sequence represents an antisense inhibitor of the invention, and inhibits the expression of the human G-alpha-13 protein. The antisense compounds of the invention are of 8 to 30 nucleobases in length, that inhibits the expression of the human G-alpha-13. The antisense compound is useful for treating an animal, particularly humans, having or being prone to a disease or condition associated with the expression of G-alpha-13, such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer; panthothenic acid; ketopanthoate reductase; panB gene; vitamin; cosmetic; medicine; nutrition; ilvC gene; panB gene; panC gene; panD gene; avtA gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Production of microorganisms that overproduce pantothenic acid, vitamin in e.g. foods or medicines, by overexpressing sequences encode ketopantothenate reductase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ņ
                                                                                                                          Antisense compound inhibiting expression of human G-alpha-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dohmen
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
3.5%; Score 14.8; DB 1; Length 18;
Best Local Similarity 88.9%; Pred. No. 1.5e+02;
Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dusch N,
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 5 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiliflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, receptor, producing proncholization, increasing levels of incr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condittion. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiatehmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sandrasagra A, Katz E,
i, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 15; SEQ ID NO 509; 872pp; English.
TACGGCATGCTGGCCCGC 140
                                                                                                                                                                                                                                                                                                                                                                                     Human oligonucleotide sequence
                                                  3 TACGGCATGATGGCCAGC 20
                                                                                                                                                                                    267/c
ABZ85267 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-APR-2002; 2002WO-US013135.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                        17-OCT-2003 (first entry)
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Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-229219/22.
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Miller S,
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The invention relates to a novel DNA vaccine for flatfish rhabdovirus (HIRRV) infected fishes, which provides immunity against HIRRV. The vaccination method uses a DNA construct comprising a transcriptional-control sequence containing cytomegalovirus immediate-type promoter, operably coupled to a nucleotide sequence encoding an immunogenic polyapeptide of HIRRV. The DNA vaccine has virucide activity. The HIRRV DNA vaccine is useful for administering to a fish belonging to the flatfish family by gene gun. The HIRRV DNA vaccine is useful for preventing HIRRV infected by HIRRV and is also useful for preventing HIRRV infection in flatfish. The HIRRV DNA vaccine is effective in enhancing immunity of fish infected by HIRRV. This effective in enhancing immunity of fish infected by HIRRV. This of polynucleotide sequence represents an oligo used in the analysis of the mRNA expression level from the muscles of flatfish, following an innoculation with the flatfish rhabdovirus vaccine of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA vaccine for flatfish rhabdovirus infected fishes has DNA construct comprising a transcriptional control sequence coupled to a nucleotide sequence encoding an immunogenic protein of flatfish rhabdovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                               DNA vaccine; flatfish rhabdovirus; HIRRV; fish; immunity;
transcriptional-control; cytomegalovirus immediate-type promoter;
immunogenic; virucide; gene gun; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.5%; Score 14.8; DB 1; Length 21; 88.9%; Pred. No. 2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 6 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Fig 5; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   54 TCAGAGGAGTCTCTGCAC 71
                                                                                                                                                                                                                                 Flatfish rhabdovirus oligo #33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MEIJ ) MEIJI SEIKA KAISHA LTD
(AOKI/) AOKI H.
CTGCACTACGAGGGCCGC 83
                              m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2001; 2001JP-00271068.
10-SEP-2001; 2001JP-00274202.
                                                                                                                                                                                                                                                                                                                                                                                                                                                26-SEP-2001; 2001JP-00294473
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                              20 CrecacrireAeeeccec
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AAA49039 standard; DNA; 20
                                                                                                                        21
                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16; Conservative
                                                                                                                        ADD22542 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 16; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-818526/77.
                                                                                                                                                                                                                                                                                                                                       Hirame rhabdovirus
                                                                                                                                                                                                                                                                                                                                                                              JP2003155254-A.
                                                                                                                                                                                            15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                27-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA49039;
                                                                                                                                                            ADD22542;
 99
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AAA49039/c
                                                                                       RESULT
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Query Match 3.5%; Score 14.8; DB 1; Length 20; Best Local Similarity 88.9%; Pred. No. 1.9e+02; Matches 16; Conservative 0; Mismatches 2; Indels

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The present invention relates to the Thermus sp. AK16D DNA ligase enzyme. This thermostable ligase has 100 fold higher fidelity than T4 ligase and 6 fold higher fidelity than Thermus thermophilus ligase. The present sequence is the degenerate antisense primer #3 corresponding to amino acids 641-647 of the T.thermophilus HB8 DNA ligase gene. This primer was used to amplify DNA ligase gene fragments from various Thermus sprains. The high specificity and thermostability of Thermus sp. Ak16D ligase hasks it useful for use in ligase based linear signal amplification, known as LDR/PCK. Ligation of suitable oligomucleotide probes can be disrupted by hybridisation mismatches. This feature may be used to detect infectious diseases (for example bacterial, fungal or viral infection),
                                                    Degenerate primer #3 targeted to T.thermophilus HB8 DNA ligase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                   New thermostable DNA ligase for sealing a ligation junction between oligonucleotide probes and the target sequence.
                                                                                     Thermostable ligase; bacterial; fungal; viral; infection; cancer genetic disease; PCR primer; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 6 C; 2 G; 5 T; 0 U; 6 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 24; 55pp; English.
                                                                                                                                                                                                                                                                                        98US-0106461P.
                                                                                                                                                                                                                                                                                                                         (CORR ) CORNELL RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic diseases and cancer
               (first entry)
                                                                                                                                                                                                                                                                                                                                                               Tong J;
                                                                                                                                           Thermus thermophilus
                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-451622/39.
                                                                                                                                                                                                                                                                                                                                                               Cao W,
                                                                                                                                                                             WO200026381-A2.
                                                                                                                                                                                                                                                     29-OCT-1999;
                                                                                                                                                                                                                                                                                        30-OCT-1998;
                 10-JAN-2001
                                                                                                                                                                                                                  11-MAY-2000,
                                                                                                                                                                                                                                                                                                                                                               Barany F,
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AAT51423-T51415 represent amplification primers used in the construction of an E. coli hygromycin B phosphotransferase (hpt) gene containing vector of the invention. The vector these sequences were used to construct also contained a luciferase gene. The hpt gene used in the vector, is used as a dominant selectable marker. The hpt gene has preferably been modified, to provide increased resistance to hygromycin in comparison to the wild type gene. In the vector, the hpt gene has the control of a promoter (such as the GDP-2 promoter) that is native to Agaricus bisporus. The vector can then be used in the production of a starker, and donor DNA are integrated into the homobasidiomycetes, and expressed at a level which allows direct selection, and stable maintenance of the transformed cells. Previoually, the donor DNA was not both integrated and expressed at high enough levels for direct selection and stable maintenance to be possible. The transformed homobasidiomycetes can be used for the commercial production of substances, such as

enzymes and metabolites

Production of stably transformed homo-basidiomycetes - with altered genetic characteristics for e.g. commercial production of enzymes.

Claim 37; Page 27; 86pp; English

Rats FH;

Huizing HJ,

Mooibroek A, Van De Rhee MD,

WPI; 1995-067335/09.

(ATOD-) ATO-DLO INST AGROTECHNOLOGISCH ONDERZOEK. (CNCC-) CNC COOEPERATIVE NEDERLANDSE CHAMPIGNONK.

94WO-NL000164.

13-JUL-1994;

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Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                      3.4%; Score 14.6; DB 1; Length 21; 81.0%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                         4; Indels
                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                             90 GACATCACCACGTCTGACCGC 110
                                                                                                                                                                                                                                                                                                                                                                                        1 GACATCACCATGGCTGAACTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human polymorphic region 313.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-US005419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0041057P.
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ26124 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 81.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (VARI-) VARIAGENICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9841648-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ26124;
                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 82
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Polymerase chain reaction; PCR; primer; amplify; E. coli; GDP-2 promoter; Agaricus bisporus; hygromycin B phosphotransferase; hpt gene; luciferase; homobasidiomycetes; metabolite; enzyme production; ss.

; 0

Gaps

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3.4%; Score 14.6; DB 1; Length 20; 63.2%; Pred. No. 2e+02; ive 6; Mismatches 1; Indels

GGCACCAAGCTGGTGAAGG 300

282 20

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12; Conservative

Local Similarity

Best Loc Matches

Query Match

BP

AAT51423 standard; cDNA; 21

RESULT

01-APR-1997 (first entry)

AAT51423;

Primer Nco-HPTS.

WO9502691-A2

Synthetic.

26-JAN-1995

Stanton VP;

Ledley FD,

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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viablity, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a precancerous condition, by administrating to the patient a first allele specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene present in a population and targets only one allelic forms of the gene comment in the disense in the disquosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, permalignant metaplastic or dysplastic concers, benigh tumours, endometriosis, polycystic kidney disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AZZSSB12-ZZSB25 represent thuman polymorphic sites described in the method of the invention
                                                                                                                     Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                               Disclosure; Fig 7; 605pp; English.
                                                                            WPI; 1998-521232/44.
                        Housman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Gaps , 0 3.4%; Score 14.6; DB 1; Length 21; 81.0%; Pred. No. 2.3e+02; ative 0; Mismatches 4; Indels Sequence 21 BP; 2 A; 12 C; 4 G; 3 T; 0 U; 0 Other; 326 GGCGGCGGACGACCAGGGCCG 346 21 gérédegakadecekédekere 1 . 0 Local Similarity 81.0 les 17; Conservative Query Match

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AAZ26557 standard; DNA; 21 BP Human polymorphic region 746. 30-NOV-1999 (first entry) AAZ26557; 

Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.

97US-0041057P 19-MAR-1998; 20-MAR-1997; WO9841648-A2 Homo sapiens 24-SEP-1998.

Stanton VP; Housman D, Ledley FD, WPI; 1998-521232/44

VARI-) VARIAGENICS INC

This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is used for preventing the development of cancer in a patient having a consciou, by administering to the patient a first allele presencerous condition, by administering to the patient a first allele present in cells of the precancerous condition, where the normal somatic consistency of the precancerous condition, where the normal somatic consistency and patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms of the gene consent in a population and targets only one allelic forms present in the ormal somatic cells, and the first gene. The products and methods can be consent in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic censions, benigh tumours, endometriosis, polygostic kindrey disease, and graft versus host disease. The method can also be used to remove malignant cells from bone marrow transplants. AAZ25812-Z26825 represent the man polymorphic sites described in the method of the invention Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease. Gaps ö Score 14.6; DB 1; Length 21; Pred. No. 2.3e+02; 0; Mismatches 4; Indels Sequence 21 BP; 2 A; 8 C; 8 G; 3 T; 0 U; 0 Other; 241 GCTGCTTCCCGGGCTCGGCCA 261 Disclosure; Fig 7; 605pp; English 21 GCGGCTTCCCAGGCAGGCCA 1 뗦. 3.4%; .042/c AAX35042 standard; DNA; 21 Query Match Best Local Similarity 81.0% RESULT 84 AAX35042/c 셤 ð

Oligonucleotide used to construct recombinant RSV vaccines. (first entry) 01-JUL-1999 AAX35042;

Respiratory syncytial virus; RSV; viral vector; mutated RSV gene; HBV; RSV antigenome; functional deletion; M2 gene; RSV-A; RSV-B; antigen; L gene mutation; vaccine; bivalent vaccine; influenza; HIV-1; HIV-2; SS. 97US-0060153P. 98US-0084133P. 98US-0089207P. 98WO-US020230 W09915631-A1 28-SEP-1998; 26-SEP-1997; 04-MAY-1998; 01-APR-1999. 12-JUN-1998; Synthetic. 

Recombinant respiratory syncytial viruses. Example 6; Page 35; 85pp; English WPI; 1999-244413/20

Bryant

Tang R, Li S,

Jin H,

(AVIR-) AVIRON INC.

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Gaps . 0

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WPI; 1999-244413/20.
                                                                                                                          (AVIR-) AVIRON INC.
                                                                                                                26-SEP-1997;
04-MAY-1998;
12-JUN-1998;
                                                                                                           28-SEP-1998;
                                                                                                 WO9915631-A1
                                                                                                      01-APR-1999
                                                                                             Synthetic.
                                                                    AAX35043;
                                                                                                                              Jin H,
                                                          RESULT 85
                                                             AAX35043
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Tang R,

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Characterizing drug-target interactions and identifying genetic mutations that confer resistance to antibacterial compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a PCR primer for the coding sequence for enoylacrate (Tabl) from Neissaria gonorrheae. The protein was used to antibacterial compounds and understand how the target and compound and understand how the target and compound and interacts. This in turn is useful for identifying other antibacterial agents. The Fabl sequence is particularly useful for generating the Hamophilus influenzae, Streptococcus pureins of N. gonorrheae, Staphylococcus aureus, Streptococcus progenes, Peculomoniae, Actinobacter, E. coli, Staphylococcus aureus, Streptococcus progenes, Pseudomonas aeruginosa, Enterococcus faecium, Bacillus subtilis and Halloobacter pylori, Which can then be used to determine how to fight infection by these bacteria. This primer was used to create random mutations in the Fabl coding sequence. (Updated on 15-SEP-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fabl; encyl-ACP reductase; DHDPE resistance; infection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.4%; Score 14.6; DB 1; Length 21; 81.0%; Pred. No. 2.38+02; ive 0; Mismatches 4; Indels
                                                           ch 3.4%; Score 14.6; DB 1; Length 21; 1 Similarity 81.0%; Pred. No. 2.3e+02; 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 5 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
           Sequence 21 BP; 3 A; 9 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neisseria gonorrheae Fabl PCR primer Gc8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 23; 55pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          266 GCACCTGGAGCAGGGGGCAC 286
                                                                                                                                                                31 GCTGGGACGAAGATGGCCACC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 dcaccrecaccarccerac 21
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                                                                                                                                                                                                                                                                                                                                                        AAA53282 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (WARN ) WARNER LAMBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neisseria gonorrhoeae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunham SA, Olson E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-350764/30.
                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200024932-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-SEP-1999;
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05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                          AAA53282;
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                                                                                                                Matches
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AAZ48457
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                                                                                                                                                                                                                   ద
The specification describes recombinant respiratory syncytial virus (RSV) particles and viral vectors which express heterologous genes or mutated RSV genes. The RSV particles comprise a RSV antigenome or genome containing at least one functional deletion in an M2 gene, or encode antigenic polypeptides of both RSV-A and RSV-B, or contain a L gene mutation. The recombinant RSV particles can be used to produce vaccines, e.g. bivalent vaccine against RSV-A and RSV-B, or RSV and influenza. Recombinant RSV vaccines can also be constructed for viruses such as HIV-BL HIV-2 and HBV, by constructing a RSV comprising a heterologous sequence from these organisms. The present oligonucleotide was used to construct the ribozyme/T7 terminator sequence, which was construct vectors which are used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The specification describes recombinant respiratory syncytial virus (RSV) particles and viral vectors which express heterologous genes or mutated RSV genes. The RSV particles comprise a RSV antigenome or genome containing at least one functional deletion in an M2 gene, or encode antigenic polypeptides of both RSV-A and RSV-B, or contain a L gene mutation. The recombinant RSV particles can be used to produce vaccines, e.g. bivalent vaccine against RSV-A and RSV-B, or RSV and influenza. Recombinant RSV vaccines can also be constructed for viruses such as HIV-1, HIV-2 and HBV, by constructing a RSV comprising a heterologous construct the ribozyme/T7 terminator sequence, which was used to vectors which are used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Respiratory syncytial virus; RSV; viral vector; mutated RSV gene; HBV; RSV antigenome; functional deletion; M2 gene; RSV-A; RSV-B; antigen; L gene mutation; vaccine; bivalent vaccine; influenza; HIV-1; HIV-2; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 3.4%; Score 14.6; DB 1; Length 21; Best Local Similarity 81.0%; Pred. No. 2.3e+02; Matches 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide used to construct recombinant RSV vaccines.
                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 2 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Recombinant respiratory syncytial viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; Page 35; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31 GCTGGGACGAAGATGGCCACC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 GCTGGGACCATGCCGGCCACC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0060153P.
98US-0084133P.
98US-0089207P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX35043 standard; DNA; 21 BP.
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hydroxylase, catecholamine-related disease, manic depression, schizophrenia, PCR primer, ss.

Rat; Nurrl; tyrosine Parkinson's disease;

Rattus norvegicus WO200058451-A1. Cell comprising exogenous nucleic acid inducing tyrosine hydroxylase expression useful for treating catecholamine-related diseases such as Parkinson's disease, manic depression and schizophrenia.

Example 3; Page 26; 68pp; English.

(SALK ) SALK INST BIOLOGICAL STUDIES.

Sakurada K, Palmer T,

WPI; 2000-656165/63.

21-MAR-2000; 2000WO-US007544

05-OCT-2000

26-MAR-1999;

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The invention provides a novel in vitro method for the detection of microorganisms and viruses. The method comprises: (1) forming a comprises and viruses. The method comprises: (1) forming a comprision of a microorganisms and viruses. The mixture by compining a predetermined volume of a sample to be tested for the presence of a nucleic acid sequence comprising 5'-GATAAGACTCCAPAAGT-3', known amounts of a first primer comprising 5'-GATAAGACTCCAPAAGT-3', known amounts of a first primer of comprising 5'-GATAAGACTCCAPAAGT-3', and a second primer comprising 5'-GATAAGACTCCTAPACTCAC-3', and primer comprising 5'-GATAGACTCTAPAGCAC-3' to replicate and attain 0.25-10000mug nucleotide product Mun mixture; (3) adding a probe containing DNA comprising 5'-GATAGACTCTTAAGCCAC-3' to the nuclet acid sequence, if present, and change the conformation of the probe; and (4) determining whether or not bacteria are present in the sample by detecting the conformational change indicating the presence of bacteria in the sample. Conformational change indicating the presence of bacteria in the sample. The methods can be used for the detection of viruses and microorganisms, including bacteria, yeast, and in clinical disapposities. Using the method for and cosmetic industry and in clinical disapposities. Using the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of microorganisms and viruses, for use in the food and cosmetic industries and for clinical diagnostics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Including bacteria, yeast, molds and protista. They can be used in food and cosmetic industry and in clinical diagnostics. Using the mult is not necessary to remove non-hybridized probe from the system
                                                                                                                                                    Microorganism, virus, polymerase chain reaction, food, cosmetic, clinical diagnostic, molecular beacon, PCR primer, ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.4%; Score 14.6; DB 1; Length 21;
81.0%; Pred. No. 2.3e+02;
iive 0; Mismatches 4; Indele
                                                                                                               Nucleic acid fragment used in detection of microorganisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 5 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rat Shh-N coding sequence PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 37; Page 38; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCTGGGACGAAGATGGCCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGTGGCTCGAAGATAGCCACC
                                                                                                                                                                                                                                                                                                                                 99WO-US010940.
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AAA95400/c
ID AAA95400 standard; DNA; 21
    21
                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                (HUNT-) HUNT WESSON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-086985/07.
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les 17; Conser
                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                     WO9963112-A2
                                                                                                                                                                                                                                                                                                                               18-MAY-1999;
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17-MAY-1999;
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                                                                          27-MAR-2000
                                                                                                                                                                                                                                                                                        09-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Romick TL,
                                     AAZ48457;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                   The present invention describes the rat Nurrl coding and protein sequences. The Nurrl protein is involved in the induction of tyrosine hydroxylase expression in adult rat-derived hippocampal progenitor cells. The Nurrl gene and protein can be used in the treatment of catecholaminerelated diseases such as Parkinson's disease, manic depression and schizophrenia. They can also be used to induce tyrosine hydroxylase expression and identify tyrosine hydroxylase related deficiencies, which are linked to the same diseases. The present sequence is a PCR primer used in a method to differentiate adult neural progenitor cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 5 A; 4 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.6; DB 1;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human gene single nucleotide polymorphism #2342.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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replace(11,a)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  230 CAAATCGGGAGGCTGCTTCCC 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 CAAATCTGACGGCTGATTCCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.4%;
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Best Local Similarity 81.0
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       X2X5X5X5X5X5X5X5X
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Mccarthy JJ;

Daley GQ,

s, Bolk

07-SEP-2000; 2000WO-US024503.

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various oblymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Infectious respiratory syncytial virus particle, useful for producing vaccines, comprises a viral genome or antigenome with a deletion in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RSV; RSV strain A2; RSV subgroup A; virus accessory gene; vaccine; ds
                                                                                                                                                                                           Nucleic acids comprising single nuclectide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide used to construct a ribozyme/T7 terminator sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ch 3.4%; Score 14.6; DB 1; Length 21; 1 Similarity 81.0%; Pred. No. 2.3e+02; 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 5 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                      (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                    Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGCTGCTCTACAGCGACTTC 365
                                                                                                                                                                                                                                                                           Example, Page 207; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bryant M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            32-AUG-2000; 2000WO-US021079.
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                               99US-0153357P.
                                        26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF25449 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                    Gargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4PI; 2001-191424/19.
                                                                                                                                                                   WPI; 2001-226749/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                              atherosclerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38-FEB-2001.
                               10-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAY-2001
                                                                                                                                     Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     345
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Matches
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The invention describes an isolated infectious respiratory syncytial varias (RSV) particle with an attenuated phenotype comprisaling an RSV antigenome or genome, where the genome or antigenome has a heterologous sequence encoding a G and P protein, and a mutation in the M2-2 gene. The SKV particle is useful as expression vector or vaccine. This sequence represents an oligonuclectide used in the construction of leader and trailer sequences for creation and functional analysis of reporter
                                                                   Oligonucleotides AAF25449-59 were used to construct a ribozyme/T7 reminator sequence, which was then ligated to the ends of the CDNA of respirators requence, which was then ligated to the ends of the constrator syncytial virus (RNV). The specification describes an infectious RNV particle comprising an RSV (anti)genome that has at least one functional deletion in a virus accessory gene. Especially, the genome contains the reverse complement of a mRNA-encoding sequence linked to a polymerase-binding site (RBS) of an RSV. The RSV particles of the invention are useful for preparing attenuated, live vaccines, including those that express heterologous gene products (particularly from another strain of RSV, some other virus or pathogen, cellular protein or tumour antigen). Also negative-strand RSV RNA templates can be used to express heterologous gene products (e.g. viral proteins or ribozymes for prevention or treatment of disease) in cells and/or to rescue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated infectious respiratory syncytial virus particle, useful as a vaccine, has an attenuated phenotype comprising the viral genome that has a heterologous sequence encoding a G and F protein and a mutation in the
                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antigenome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Respiratory syncytial virus genome construction oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Respiratory syncytial virus; RSV; attenuated phenotype; antigenoi
G protein; F protein; M2-2 gene; expression vector; vaccine; ss.
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                                                                                                                                                                                                                                                                                                                                                                                             Score 14.6; DB 1; Length 21;
Pred, No. 2.3e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                             Seguence 21 BP; 2 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                    Disclosure; Page 34; 128pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 39; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31 GCTGGGACGAAGATGGCCACC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-NOV-2000; 2000US-00724416.
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Best Local Similarity 81.0%;
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ď
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M2-2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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ABK96224/c
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Gaps

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1 CTGATTGACAGGGACTTCCTC 21

BP

21

ADC49462 standard;

(first entry)

18-DEC-2003

ADC49462;

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ADC49462
ID ADC4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a drug composition containing as an active component a substance which activates FC receptor gamma chain. Also included are detecting oligodendroglia and their precursor cells capable of forming myelin (using as an indicator the expression of FC receptor gamma chain in the precursor cells), investigation of the expression of FC receptor gamma chain in the precursor cells), investigation of the expression of FC receptor gamma chain in animal brain tissue (by immune typing, cytochemical amalysis, gene amplification or Western blotting) and kits for these methods, containing anit-FC receptor gamma chain antibody or amplification primers. The drug is used for treatment and prevention of neurodegenerative diseases, and disorders of myelin formation, such as multiple solerosis, Krabbe's diseases, advenoledystrophy and multiple solerosis, Krabbe's diseases, advenolethodystrophy and metachromic leukodystrophy. The present sequence is an FC receptor III alpha gamma chain PCR primer used in the exemplification of the invention
plasmids and construction of a cDNA representing the complete genome of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR; primer; ss; Fc receptor gamma chain; neuroprotective; oligodendroglia; myelin; neurodegenerative disease; unlitiple sclerosis; myelin formation disorder; Krabbe's a disease; adrenoleukodystrophy; metachromic leukodystrophy; Fc receptor III alpha gamma chain.
                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Drug compositions containing Fc receptor gamma chain activator for treatment and prevention of neurodegenerative disorders including multiple sclerosis.
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                                                                                           / Match 3.4%; Score 14.6; DB 1; Length 21; Local Similarity 81.0%; Pred. No. 2.3e+02; les 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 21 BP; 4 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
                                                        Seguence 21 BP; 2 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fc receptor III alpha gamma chain PCR primer #2.
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                                                                                                                                                                                 31 GCTGGGACGAAGATGGCCACC 51
                                                                                                                                                                                                             21 GCTGGGACCATGCCGGCCACC 1
                                                                                                                                                                                                                                                                                                                             ABX95654 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
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                                                                                                                                                                                                                                                                                                                                                                    ABX95654;
                                                                                              Query Match
                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                     RESULT 92
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                                                                                                      non-human animal model; demyelinating disease; myelinogenesis inhibition; myelinogenesis signal molecules; oligodendroglia; screening; myelin growth regulator; multiple sclerosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises a non-human animal model for demyelinating disease - in which myelinogenesis is inhibited by a defect of myelinogenesis signal molecules in oligodendrogila. The non-human animal model of the invention is useful for acreening for a myelin growth regulator, or for screening for a therapeutic agent which is useful for treating a demyelinating disease such as multiple sclerosis. The present DNA sequence represents a PCR primer that was used in an example of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Multipotent adult stem cell; MASC; cell replacement therapy; cytostatic; cardiant; cardiovascular; hepatotropic; haemostatic; antidiabetic; virucide; antiinflammatory; vasotropic; antianaemic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                  Novel non-human animal model for demyelinating disease in which myelinogenesis is inhibited by defect of myelinogenesis signal molecules in oligodendroglia, for screening for therapeutic agent for multiple
                                                                             Non-human animal model for demyelinating disease-related PCR primer #10.
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Local Similarity 81.0%; Pred. No. 2.3e+02;
Les 17; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 4 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                       (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; SEQ ID NO 10; 56pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              348 CTGCTCTACAGCGACTTCCTC 368
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                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-630032/60.
                                                                                                                                                                                                            JP2003079270-A.
                                                                                                                                                                           Unidentified.
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25-NOV-2002
                                                                                                                                                                                                                                           18-MAR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
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348 CIGCICTACAGCGACTICCIC 368

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Retrovirus LTR PCR primer.

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New multipotent adult stem cells that can be induced to differentiate to form a cell type of mesodermal, ectodermal or endodermal origin, useful for treating e.g. cancer, diabetes, hepatitis, hemophilia, ischemia or
cerebroprotective; immunosuppressive; antibacterial; PCR; primer; ss.
                                                                                                                                                                                                          Example 10; Page 55; 117pp; English.
                                                                                       14-FEB-2001; 2001US-0268786P.
15-FEB-2001; 2001US-026962P.
7-AUG-2001; 2001US-0343386P.
25-OCT-2001; 2001US-0343386P.
                                                                        14-FEB-2002; 2002WO-US004652
                unidentified retrovirus.
Unidentified.
                                                                                                                                                 WPI; 2002-667000/71.
                                                                                                                                 (ANON ) ANONYMOUS.
                                        WO200264748-A2.
                                                                                                                                                                                          inflammation.
                                                       22-AUG-2002.
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The present sequence is a primer for a retrovirus long terminal repeat

(LTR). The primer was used in an example from the invention in which a
retroviral marking study was used to demonstrate that neurons, astrocytes
and oligodendrocytes can produced from a single bone marrow-derived
multipotent adult stem (MASC), which also differentiated into muscle and
addifferentiating MASC. The MASC are derived from a non-embryonic organ
or tissue, such as bone marrow, muscle, brain, umbilical cord blood or
placenta, and have the capacity to be induced to differentiate to a cell
type of mesodermal, ectodermal or endodermal origin, including
of mesolast, chondrocyte, adipocyte, fibroblast, marrow stroma, skeletal
muscle, smooth muscle, cardiac muscle, endothelial, pitter,
coffeeblast, chondrocyte, adipocyte, fibroblast, marrow stroma, skeletal
muscle, smooth muscle, cardiac muscle, endothelial, pitter,
coffeeblast, chondrocyte actian muscle, endothelial, spittelial, liver,
muscle, smooth muscle, cardiac muscle, endothelial, pitter,
coffeeblast, constitutively express oct4 and high levels of telomerase and are
negative for CD44, MHC class I and MHC class II expression. Teratomas are
not formed when MASC are administered to a patient. MASC or their progeny
are particularly useful for treating cancer, cardiacyascular disease,
coffeciency connective tissue disorders, autoimmuse disease,
degenerative or traumatic neurological conditions, autoimmuse disease,
degenerative or traumatic neurological conditions, autoimmuse disease,
pe directed to abdominal aortic aneurysm, cardiac bypass surgery,
peripheral vascular disease, or coronary vascular disease (all claimed).
(Updated on 07-AUG-2003 to correct OS field) (Updated on 29-AUG-2003 to
standardise OS field)

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Gaps
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                                                      Query Match 3.4%; Score 14.6; DB 1; Length 22; Best Local Similarity 81.0%; Pred. No. 2.5e+02; Matches 17; Conservative 0; Mismatches 4; Indels
Sequence 22 BP; 3 A; 4 C; 8 G; 7 T; 0 U; 0 Other;
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(revised)
(revised)
(first entry) 29-AUG-2003 07-AUG-2003 25-NOV-2002 ABN84966; RESULT 95
ABN84966/C
ID ABN84
XX
AC ABN84
DT 29-AU
DT 07-AU
DT 25-NO

.966/c ABN84966 standard; DNA; 22

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Multipotent adult stem cell; MASC; cell replacement therapy; cytostatic; cardiant; cardiovascular; hepatotropic; haemostatic; antidiabetic; virucide; antiinflammatory; vasotropic; antianaemic; neuroprotective; cerebroprotective; immunosuppressive; antibacterial; PCR; primer; ss.
                                                                                                                14-FEB-2001; 2001US-0268786P.
15-FEB-2001; 2001US-0269062P.
07-AUG-2001; 2001US-0310625P.
25-OCT-2001; 2001US-0343386P.
                                                                                                    14-FEB-2002; 2002WO-US004652
                                                     unidentified retrovirus.
Unidentified.
                                                                                                                                                    (ANON ) ANONYMOUS.
                                                                          WO200264748-A2
                                                                                       22-AUG-2002.
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New multipotent adult stem cells that can be induced to differentiate to form a cell type of mesodermal, ectodermal or endodermal origin, useful for treating e.g. cancer, diabetes, hepatitis, hemophilia, ischemia or inflammation.

WPI; 2002-667000/71.

Example 10; Page 55; 117pp; English.

The present sequence is a primer for a retrovirus long terminal repeat (ITR). The primer was used in an example from the invention in which a retroviral marking study was used to demonstrate that neurons, astrocytes and oligodendrocytes can produced from a single bone marrow-derived multipotent adult stem (MASC), which also differentiated into muscle and differentiating masc, made marrow, muscle, brain, umbilical cord blood or placenta, and have the capacity to be induced to differentiate to a cell type of mesodermal, ectodermal or endockmal origin, including or tissue, such as bone marrow, muscle, brain, umbilical cord blood or placenta, and have the capacity to be induced to differentiate to a cell type of mesodermal, ectodermal or endockmal origin, including coteoblast, chondrocyte, adipocyte, fibroblast, marrow stroma, skeletal coteoblast, smooth muscle, cardiac muscle, endothelial, polithelial, liver, muscle, smooth muscle, cardiac muscle, endothelial, epithelial, liver, cot panoreas, haematopoletic, gllal, neuronal or oligodendrocyte cell types. MASC constitutively express oct4 and high levels of telomerase and are negative for CD44, MHC class I and MHC class II expression. Teratomas are cot farmed when MASC are administered to a partient. MASC or their progeny are particularly useful for trreating cancer, cardiovascular disease, car eadministered to a particinal conditions, autoimmune disease, desencative or traumatic neurological conditions, autoimmune disease, desence, transplant rejection, ischaemia or inflammation. Treatment may be directed to abdominal aortic aneurysm, cardiac bypass surgery, peripheral vascular disease, or coronary vascular disease (all claimed). (Updated on 07-MUG-2003 to correct OS field.) (Updated on 29-AUG-2003 to cytandardise OS field)

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Gaps
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                                          Query Match
3.4%; Score 14.6; DB 1; Length 22;
Best Local Similarity 81.0%; Pred. No. 2.5e+02;
Matches 17; Conservative 0; Mismatches 4; Indels
Sequence 22 BP; 3 A; 4 C; 8 G; 7 T; 0 U; 0 Other;
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21 Arcacrcadaddadadccrcc 1 RESULT 96 AAC72998/c ID AAC72998 standard; DNA; 17 BP. ద

09-FEB-2001

AAC72998;

WO200058519-A2. Homo sapiens.

05-CCT-2000.

31-MAR-1999;

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Altshuler D, Cargill M,
Lipshutz RJ, Patil N, 6
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                                                                                                                                                                                                                                                                                                                          genetic analysis.
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Lipshutz RJ,
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                                                                                                 31-MAR-1999;
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                               05-OCT-2000.
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                                                                                                            Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis.
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Lipshutz RJ, Patil N, Sklar P;
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                                                                           Single nucleotide polymorphism PCR primer #1885.
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                                            (first entry)
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09-FEB-2001

AAC72992;

AAC72992/c RESULT 97

Query Match Best Local S Matches 15

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diseases

Homo sapiens

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The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in diesase diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose usceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, echizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
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                                                                                                                                                                                                                                                                                   , Daley GQ, Ireland JS, Lander ES;
Sklar P;
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                                                                                                                                                             (WHED ) WHITEHEAD INST BIOMEDICAL RES. (APFY-) APPYMETRIX INC.
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30-MAR-2000; 2000WO-US008440.
                                                                                  99US-0127248P
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04-DEC-2000
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                                                                                                                                                                     The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an
                                                     Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lander
                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , Daley GQ, Ireland JS, Sklar P;
                                                                                                                                                                                                                                                                                                                                                        Seguence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                    3.4%; Score 14.4; DB 1;
93.8%; Pred. No. 1.6e+02;
iive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Single nucleotide polymorphism PCR primer #1889.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (WHED ) WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 8; Fig 5; 214pp; English.
                                                                                                                                             Claim 8; Fig 5; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGAGCCCCGGGGACC 318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTGAGACCCGGGGGACC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAC73004/c
ID AAC73004 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Patil N, S
                                                                                                                                                                                                                                                                                                                                                                                                                             15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-611722/58.
                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                             genetic analysis.
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Lipshutz RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-MAR-1999;
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individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNBs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
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                                                                                                                                                                                                                                                        3.4%; Score 14.4; DB 1; Length 17; 93.8%; Pred. No. 1.6e+02; Ative 0; Mismatches 1; Indels
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                                                                                                                                                                                                       Sequence 17 BP; 1 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 60; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Barber JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cdk8 ribozyme binding site #90.
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                                                                                                                                                                                                                                                                                                                                                                                  303 CTGAGCCCCGGGGACC 318
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tes 15; Conservative
                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-412314/35.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-DEC-1998;
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dopaminergic gene; DRD5; susceptibility diagnosis; h aura; depression; anxiety; variant allele detection;

migraine with aura; differentiation; ss

PCR primer;

Homo sapiens. WO9807426-A1.

Synthetic

97WO-US014830. 96US-0024399P. 97US-0036091P.

21-AUG-1997; 22-AUG-1996; 17-JAN-1997;

26-FEB-1998

(GLAX ) GLAXO GROUP LTD

Peroutka SJ;

Primer 40DRD5.SB.PCR2 for DRD5 gene.

(first entry)

09-JUL-1998

AAV25487;

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AAV25487 standard; DNA;

AAV25487

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule [11] comprising a promoter operably linked to a nucleic acid segment encoding [1]. [1] can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, chanded and segment encoding [1]. [1] can have antipsoriatic, and percention optical, wulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. [1] can be used in gene therapy. [1] and [1] are useful for treating proliferative skin diseases such as periasis, atopic dermatitis, actinic keratosis, also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing carring such as keloid, adhesion and whypertrophic or hypertrophic burn scar. AMH57577 to AMH62099 represent sequences used in the companies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; WNE; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                        Cell-cycle dependent kinase cdk8 ribozyme binding site SEQ ID NO:956.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 4 A; 3 C; 9 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-OCT-2000; 2000WO-US029500
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AAH58532
ID AAH58532 standard; DNA; 19
                                                                                                                                                                                                                                          10-SEP-2001 (first entry)
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is a primer for the dopaminergic gene DRD5, which can be used in a novel method assess susceptibility to a syndrome having symptoms of migraine with aura (MMA), depression and/or anxiety. The method comprises detecting a variant allele of at least 1 dopaminergic gene. Analysis of variant dopaminergic gene alleles may also differentiate between patients with MMA, and those with migraine without aura (MO) or other conditions, e.g. stroke, that produce similar symptoms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:127.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Assessing susceptibility to syndromes including migraine with aura, depression and anxiety - by detecting variant alleles in genes for dopaminergic receptors or transporter, also treatment using agents that antagonise binding of dopamine to these proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EPB2; LAP, TCF5; CRP2; NFIL6; IL6DBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsiveness;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 7 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.4%; Score 14.4; DB 1;
93.8%; Pred. No. 2.2e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 18; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  204 GTGAAAGCAGAGACT 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABL94361 standard; DNA; 20 BP.
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Best Local Similarity 93.8'
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-168887/15.
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Gaps

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3.4%; Score 14.4; DB 1; Length 19; 93.8%; Pred. No. 2e+02; ive 0; Mismatches 1; Indels

GGTGAAGGACCTGAGC 308

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Local Similarity nes 15; Conserv

Query Match Best Loc Matches 16

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Gaps . 0

Indels

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Mismatches

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15; Conservative

Matches

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oxidative stress response; IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothicate; antisense oligonucleotide; ss.
                                        Key
modified_base
                                                                                      modified base
                                                               modified base
                               Mus musculus
                                                                                                                  US6271030-B1
                                                                                                                           07-AUG-2001
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/mod\_base= OTHER /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE cytosines are 5-methylcytosine"

/mod\_base= OTHER /note= "Phosphorothioate linkages"

Location/Qualifiers

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1. .20 /\*tag=

// Caracas OTHER // (2'-MOE) nucleotides. All 2' MOE cytosines are 5-methylcytosine"

U 16. .20

/\*tag=

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Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BBP alpha) and which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human and/or mouse C/BBP alpha RNA, and were analysed for their effect on C/BBP alpha mENA levels of transcription factors which regulate the expression of a wide range of transcription factors which requiate the expression of a wide range of contranscription factors which requiate the expression of a wide range of contranscription factors which requiate the expression of a wide range of configuratily regulates hormone responsiveness and oxidative stress responses of primarily regulates hormone responsiveness and oxidative stress responses of primarily regulates hormone responsiveness and oxidative stress responses and is a mediator of IL-6 (interleukin-6) signalling, C/BBP beta is thought to be involved in carbohydrate metabolism, immunity, the Thi response, female ferrility and gluconeogenic pathways. C/BBP beta is thoughts to be involved in the liver, lung, spleen, kidney, brain, and testils, with the highest expression found in the lung. It is also expressed at a higher of elevated levels of glucose, indicating that it is involved in the conditions associated with C/BP beta expression, conditions are useful for diagnosis, prevention and treatment of manimum cancer), tunour formation, diabetes (particularly covarian cancer), tunour formation, diabetes (particularly covarian cancer), tunour formation, diabetes (particularly covarian cancer), tunour formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/EBP) beta, useful in vitro for inhibiting expression of human or mouse C/EBP beta in cells/tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Col 47-48; 69pp; English
14-JUN-2000; 2000US-00593711.
                                                                                                          14-JUN-2000; 2000US-00593711.
                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-214451/27.
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3.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 2.2e+02;
Sequence 20 BP; 1 A; 6 C; 8 G; 5 T; 0 U; 0 Other;
                                                       Query Match
Best Local Similarity
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Sequences ABL94252-ABL94476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BSP alpha) gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human and/or mouse C/EBP alpha RNA, and were analysed for their effect on C/BBP alpha mRNA levels by quantitative real-time PCR. The C/EBP family of proteins are a family of transcription factors which regulate the expression of a wide range of genes that control normal tissue development, cellular function, cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= c/mod_base OTHER
/mod_base OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE
cytosines are 5-methylcytosine"
                                                                                                                                                                                                       Mouse C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:128.
                                                                                                                                                                                                                              Mouse; murine; C/EBP beta; CCAAT/enhancer-binding protein beta; C/EPB2; LAP, TOP5; CRP2; NFIL6; IL6DBP; NF-M; AGP/EBP; Apc/EBP; transcription factor; tissue development; cellular function; proliferation; differentiation; hormone responsivenes; cardative stress response; IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; esc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2'
cytosines are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/EBP) beta, useful in vitro finhibiting expression of human or mouse C/EBP beta in cells/tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
1. 20
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Col 49-50; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= b
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUN-2000; 2000US-00593711.
383 CGACGACGCCCCAAG 398
                                                                                                               ABL94362 standard; DNA; 20
                                                                                                                                                                            29-JUL-2002 (first entry)
                       16 CGACTACGGCGCCAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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                                                                                                                                               ABL94362;
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proliferation and functional differentiation. C/EBP beta (also known as C/EPB2, LAP, TCF5, CRP2, NFIL6, IL6DBP, NF-M, AGP/EBP and Apc/EBP)

primarily regulates hormone responsiveness and oxidative stress responses and is a mediator of IL-6 (interleukin-6) signalling. C/EBP beta is thought to be involved in carbohydrate metabolism, immunity, the Thi response, female fertility and gluconeogenic pathways. C/EBP beta is capponse, female fertility and gluconeogenic pathways. C/EBP beta is response, female fertility and gluconeogenic pathways. C/EBP beta is nighest expression found in the lung. It is also expressed at a higher level in malignant ovarian tissue compared with normal ovarian tissue, and its expression in pancreas is upregulated in response to chronically elevated levels of glucose, indicating that it is involved in the impairment of insulin secretion in type II diabetes. The oligonucleotides of the invention are useful for diagnosis, prevention and treatment of conditions associated with C/EBP beta expression, such as cancer (particularly ovarian cancer), tumour formation, diabetes (particularly type II diabetes), infection, or inflammation 888888888888888888888888

Sequence 20 BP; 1 A; 6 C; 8 G; 5 T; 0 U; 0 Other;

ö 3.4%; Score 14.4; DB 1; Length 20; 93.8%; Pred. No. 2.2e+02; ive 0; Mismatches 1; Indels 383 CGACGACGCCCCAAG 398 Ŋ CGACTACGGCGCCAAG Best Local Similarity 93.8 Matches 15; Conservative 20 ઠે a

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RESULT 105 AAV32912/c ID AAV32912 standard; DNA; 21 AAV32912; Bovine lactoferrin cDNA primer 1.

(first entry)

26-OCT-1998

PCR; primer; amplification; pepsin; gastrointestinal tract; milk; Aspergillus niger beta-galactosidase gene; lactase intolerance; cheese making; chymosin; bovine lactoferrin cDNA; ss.

Synthetic,

WO9829536-A2

09-JUL-1998

29-DEC-1997;

96US-00775842. 31-DEC-1996;

97WO-IB001658

(NEXI-) NEXIA BIOTECHNOLOGIES INC.

Kabel JJ, Amantea GF Turner JD, Eino M, Karatzas CN,

Mccarthy JJ

Bolk S, Daley GQ,

Gargill M, Ireland JS,

Lander ES,

WPI; 2001-226749/23

Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and

Example, Page 48; 242pp; English.

atherosclerosis.

WPI; 1998-388118/33.

- can be useful Synthetic beta-galactosidase inactive in milk but active in vivo chemically activated and used to treat lactose intolexance, also in cheese production.

Example 1; Page 13; 38pp; English.

Primers 1 and 2 (AAV32913) were used in a PCR reaction to amplify the bovine lactoferrin cDNA. The PCR product was used as a tail which was used through a pepsin recognition site to the 3' end of the Aspergillus niger beta-galactosidase gene. The invention provides a synthetic betagalactosidase which differs from the natural occurring enzyme in being classified in milk but capable of being activated by a chemical or condition naturally present in the gastrointestinal tract of humans. The design of this synthetic enzyme comprises of a tail domain fused to the

The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various oblymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also

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beta-galactosidase through a cleavage site. The presence of the tail domain renders the enzyme inactive and it can also be used as a purification handle. The synthetic beta-galactosidase is claimed to be able to hydrolyse lactose in vivo to overcome lactase intolerance and thereby reduce associated gastrointestinal disorders. The synthetic beta-galactosidase is also claimed to be useful in cheese making whereby it is activated by chymosin when added to milk
                                                                                                                                                                                                                                                                                                                                                                                                  Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; cocronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                    Query Match 3.4%; Score 14.4; DB 1; Length 21; Best Local Similarity 93.8%; Pred. No. 2.5e+02; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                           Sequence 21 BP; 4 A; 5 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                         Human gene single nucleotide polymorphism #16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
replace(11,G)
/*tag= a
                                                                                                                                                                                                                                                                                         AAF95255 standard; DNA; 21 BP.
                                                                                                                                                                                           192 ATCCACTGCTCGGTGA 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-SEP-2000; 2000WO-US024503
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                                                                                                                                                                                                                      18 Arccagracrogram
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                                                                                                                                                                                                                                                                                                                                               06-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-MAR-2001.
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Variation
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Sequence 21 BP; 6 A; 6 C; 8 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                  Human, variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
    useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bolk S, Daley GQ, Mccarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/standard_name= "single nucleotide polymorphism"
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                                                                         3.4%; Score 14.4; DB 1; Length 21; 93.8%; Pred. No. 2.5e+02; tive 0; Mismatches 1; Indels
                                                    Sequence 21 BP; 7 A; 5 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                             Human gene single nucleotide polymorphism #1169.
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                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers replace(11,T)
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                                                                                                                              204 GIGAAAGCAGAGACT 219
                                                                                                                                                                                                                   AAF96408 standard; DNA; 21 BP
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2000US-0225724P
                                                                                                                                                      GTGAATGCAGAGAACT 21
                                                                                                                                                                                                                                                                      06-JUN-2001 (first entry)
                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-226749/23
                                                                                       Best Local Similarity Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  atherosclerosis.
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26-JUL-2000;
16-AUG-2000; 2
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                                                                                                                                                                                                                                                                                                                                                                                                              Key
Variation
                                                                                                                                                                                                                                              AAF96408;
                                                                               Query Match
                                                                                                                                                                                            RESULT 107
                                                                                                                                                                                                           AAF9640E
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                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                         yak milk; alpha-lactoalbumin; beta-lactoglobulin; alpha S1-casein; alpha S2-casein; beta-casein; kappa-casein; lactoferritin; ss.
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Score 14.4; DB 1; Length 21; Pred. No. 2.5e+02;
                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 6 A; 4 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 5 (disclosure); 41pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seven kinds of yak milk protein gene sequence.
                                                 Mismatches
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                                                                                              301 ACCTGAGCCCCGGGGA 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28 AGGCTGGGACGAAGA 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-DEC-2000; 2000CN-00134189
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    3.4%;
Local Similarity 93.8%;
les 15; Conservative (
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AAT13226 standard; DNA; 22
                                                                                                                                                                                                                                                            ADE64663 standard; DNA; 21
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nes 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-741796/81.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bos grunniens.
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                                                                                                                                                                                                                                                                                                                                                     29-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JUL-2002.
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         Query Match
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WPI; 1996-174574/18.
                                                    (TOYJ ) TOSOH CORP.
                                                                  concurrently in in supernatant.
                                                                                                                                                                                                    Homo sapiens
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                                        11-AUG-1994;
                                              11-AUG-1994;
                            JP08051995-A
                                                                                                                                                                                                               11-JJN-1998
                                  27-FEB-1996.
                                                                                                                                                                                                 Synthetic
                      Synthetic.
                                                                                                                                                             AAV42250;
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                                                                                                                        Query Match
                                                                                                                                                  RESULT 110
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Disclosure, Page 30, 128pp; English. Human universal VH primer MG-30. . 0 Jakobovits A, Kucherlapati R, AAV68617 standard; DNA; 22 BP. 98WO-US009160. 263 GGTGCACCTGGAGCAG 278 3.4%; GGTGCAGCTGGAGCAG 18 97WO-US023091 96US-00759620 (first entry) Query Match Best Local Similarity 93.8' Matches 15; Conservative WPI; 1998-33314/29. (ABGE-) ABGENIX INC. of human antibodies 05-MAY-1998; Synthetic. Homo sapiens WO9850433-A2 03-DEC-1997; 03-DEC-1996; 30-MAR-1999 12-NOV-1998. AAV68617; m 111 ద #X#X#X#X#####X#X####X#X ઠ Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa; production; antibody; PCR primer; 88. The present sequence is a PCR primer for plasmid pBlue-TH6, which was used in the construction of the expression plasmid pEdHCkappa-TH6. pEdHCkappa-TH6 was prepd. by inserting a gene encoding the light-chain (LC) variable region of human anti-TSH antibody (Ab) into pEdHCkappa, an expression vector for an Ab LC. pEdHCkappa-TH6, an expression vector for the prodn. of an Ab LC in an animal host cell, contains 5'3' a SV40 promoter, and base sequences encoding dihydrofolate reductase, Ab LC signal sequence and Ab LC variable and constant regions. pEdHCkappa-TH6 along with the equivalent heavy chain expression vector pEdHCG1-TH8 can be used for the prepn. of an Ab mol. in an animal host cell Expression vectors for antibody (Ab) heavy and light chains - introduced concurrently into animal host cell to produce Ab mol. which is secreted Gaps Plasmid; pBlue-TH6; construction; expression plasmid; primer; pBdHKappea-TH6; light chain; variable region; human; antibody; TSH; thyroid stimulating hormone; animal host cell; SV40 promoter; dihydrofolate reductase; PCR; polymerase chain reaction; ss. ; 0 3.4%; Score 14.4; DB 1; Length 22; 93.8%; Pred. No. 2.7e+02; tive 0; Mismatches 1; Indels Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other; Plasmid pBlue-TH6 PCR primer TSEVK1FOR Universal human VH PCR primer MG-30. Example 2; Page 7; 9pp; Japanese 297 BP 94JP-00189277 94JP-00189277 7 AAV42250 standard; DNA; 22 23-SEP-1998 (first entry) GGCACCAAGCTGGTGA 22 égcaccaagerregaga Local Similarity 93.8 nes 15; Conservative

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cranscripts expressed in Xenomice. The products were used for repertoire transcripts expressed in Xenomice. The products were used for repertoire analysis. The specification describes a transgenic non-thuman mammal which has genome modification describes a transgenic non-thuman mammal which cours, so that the mammal does not display normal B-cell development. The could genome also has an inserted human heavy chain Ig locus in germline configuration, the human heavy chain Ig locus in micro constant region and regulatory and switch sequences, human J-H genes and an inserted human J-H micro constant region and regulatory and switch sequences, human J-H genes, and human V-H genes and an inserted human J-H collight chain Ig locus in germline configuration, the human kappa light chain Ig locus in mermine configuration, the human kappa light chain Ig locus where the number of V-H and V-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The cransgenic animals have a near complete human Ig locus, including both a transgenic animals have a near complete human in locus. They can be used for the production of human antibodies when exposed to particular anigens e.g. when exposed to human antibodies when respectively
                                                                                                                New transgenic non-human mammals - having an inactivated immunoglobulin
locus and a near complete human immunoglobulin locus, used for production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; PCR; primer; amplification; human; epidermal growth factor receptor; tumour; BGF; transforming growth factor alpha; TGF-alpha.
Green L;
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Pred. No. 2.7e+02;
0; Mismatches 1; Indels
   Mendez M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
   Klapholz S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jia X;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gallo M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yang X,
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The primers AAV68617-V68618 were used to produce anti-epidermal growth factor receptor (EGF-r)-antibodies. The antibodies can be administered therapeutically to patients (human or veterinary) to treat solid tumours EGF-r is overexpressed on many human solid tumour types, and the fully human antibodies (i.e. comprising (i) and (ii)) inhibit both epidermal growth factor (EGF) and transforming growth factor alpha (TGF-alpha) binding to EGF-r (Known to lead to cellular proliferation and tumour growth). They can prevent tumour cell growth and, in combination with an antineoplastic agent, may eradicate established tumours. The fully human antibodies can minimise the immunogenic and allergic responses intrinsic to previous mouse/rat or mouse/rat-derived antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sox-9; bone regeneration; cartilage regeneration; campomelic dysplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated SOX-9 genes - used to develop prods. for the promotion or suppression of bone or cartilage differentiation of growth.
                               Humanised antibodies against epidermal growth factor receptor, EGF-r useful to treat solid tumours whilst inducing reduced immunogenic or allergic effects compared to mouse or mouse-derived antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                3.4%; Score 14.4; DB 1; Length 22; 93.8%; Pred. No. 2.7e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy; sex reversal; primer; single strand conformation polymorphism; SSCP; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22 BP; 4 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 42; 64pp; English.
                                                                                                                Example 3; Page 96; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     263 GGTGCACCTGGAGCAG 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-NOV-1994; 94AU-00009714.
05-DEC-1994; 94AU-0009835.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95WO-AU000799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGTGCAGCTGGAGCAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT30310 standard; cDNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Goodfellow PN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                  3.47
Best Local Similarity 93.89
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYQU ) UNIV QUEENSLAND. (UYCA-) UNIV CAMBRIDGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOX-9 SSCP primer 534.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-277777/28.
WPI; 1999-034712/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9617057-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Koopman PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .06-JUN-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT30310;
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AAT3031
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SSCP primers 534 (AAT30310), 661 (AAT30311), 687 (AAT30312), 854 (AAT30313), 836 (AAT30314) and 1018 (AAT30315) were used for SSCP analysis of the SOX-9 gene (see also AAT30309) in campomelic dysplasia

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Gaps ö

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                                                                                                                                                             ö
(CD) patients. Primers were designed to amplify the known coding sequence and intron/exon junctions. Unique SSCP conformers were cloned and sequenced. Alterations in SOX-9 can cause both CD and male-to-female sex reversal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human glial cell derived neurotrophic factor and its derivatives and use.
                                                                                                                                                                                                                                                                                                                                                                                                                                 derived neurotrophic factor; GDNF; PCR; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                    Human glial cell derived neurotrophic factor (GDNF) PCR primer #44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                              Length 19;
                                                                                                                                                             3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 3 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                              Sequence 19 BP; 7 A; 2 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                           7 Match 3.3%; Score 14.2; DB 1; Local Similarity 94.2%; Pred. No. 2.2e+02; les 16; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (YISH-) YISHENG BIOLOGICAL PHARM CO LTD SHUHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 6; Page 4 (Claims); 28pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         228 GCCAAATCGGGAGGCTGCT 246
                                                                                                                                                                                              350 GCTCTACAGCGACTTCCTC 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 GCGGAATCGGCAGGCTGCT 1
                                                                                                                                                                                                                    19 GTTCTTCACCGACTTCCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ73805 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                             ACA96850 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-JAN-2001; 2001CN-00107450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JAN-2001; 2001CN-00107450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhou S, Zheng Z, Feng H;
                                                                                                                                                                                                                                                                                                                                                                          24-JUL-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             nervous system disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-000523/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, glial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CN1364812-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-AUG-2002.
                                                                                                                                   Query Match
Best Local Si
Matches 16;
                                                                                                                                                                                                                                                                                                                                               ACA96850;
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AAQ73805/c
ID AAQ7380
XX
                                                                                                                                                                                                                                                                                               ACA96850,
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(revised) (first entry)

25-MAR-2003 22-MAY-1995

AAQ73805;

Aspergillus aculeatus; ss

Pectin lyase,

Aspergillus aculeatus

WO9421786-A1

29-SEP-1994

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which are complementary to at least a portion of the human osteopontin (OPN) cDNA sequence (AAX29191). The antisense sequences are used to prevent restenosis in tissue, particularly coronary arterial tissue, especially where the patient is undergoing angioplasty, particularly percutaneous trans-luminal coronary angioplasty or directional coronary atherectomy. They prevent secretion of osteopontin by monocytes and macrophages which infiltrate to sites of inflammation following surgery osteopontin probably causes restenosis by inducing coronary artery smooth muscle cells (CASMC) to migrate to, and proliferate at, angioplasty injury sites. Sequences AAX2917-178 represent RT-PCR primers specific
                                                                                                                                                                                                                                                                                                               relates to antisense osteopontin oligonucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, mitochondrial phosphoenolpyruvate carboxykinase; PBPCK-M; PCK2; PEPCK-mitochondrial; mtPEPCK, antisense oligonucleotide; modulation; phosphorothioate; inhibition; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mtPEPCK phosphorothioate antisense oligonucleotide SEQ ID NO:27
                                                                                                                                                                                                                  New osteopontin antisense sequences - useful to treat restenosis, particularly following vascular surgery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.2; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                           & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for human osteopontin cDNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                Panda DK;
                                                                                                                                                                                                                                                                             Example 1; Page 28; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          48 CACCACTCAGAGGAGTCTC 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 caccagrerdardadrere 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.3%;
                                   98WO-US016569
                                                                       97US-0054967P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ95339 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16; Conservative
                                                                                                                                                Kundu GC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
                                                                                                           (USSH ) US DEPT HEALTH
                                                                                                                                                                                    WPI; 1999-190049/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
Synthetic.
                                                                                                                                                  Mukherjee AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-AUG-1999;
                                                                         07-AUG-1997;
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                                   07-AUG-1998;
18-FEB-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ95339;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 116
AAZ95339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ73789-Q73822 are partial DNA sequences, one or more of which can be used to encode enzymes having pectin lyase (PL) activity. The Aspergillus sequence by the product the separatial sequences were derived are shown in AAR60881 and AAQ73823 respectively. These PL enzymes degrade plant cell wall components, and can therefore be used to reduce the viscosity of fruit juices. They can also be used for the production of antibodies. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Osteopontin, antisense, restenosis, coronary arterial tissue; CASMC; inflammation, coronary artery smooth muscle cell; angioplasty; human; OPN; RT-FCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                cell wall degradation; reducing fruit juice viscosity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New pectin lyase enzyme from Aspergillus aculeatus - used for the degradation of plant cell wall components, esp. for reducing the viscosity of fruit juices.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Christgau S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.3%; Score 14.2; DB 1; Length 20;
84.2%; Pred. No. 2.4e+02;
ive 0; Mismatches 3; Indels
                                                                                             Aspergillus aculeatus pectin lyase partial DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human osteopontin (OPN) specific RT-PCR primer hOPN-L.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Kofod LV, Kauppinen MS, Andersen LN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other;
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Claim 2; Page 47; 65pp; English.

94WO-DK000105

11-MAR-1994;

93DK-00001216

12-MAR-1993; 28-OCT-1993;

(NOVO ) NOVO-NORDISK AS

WPI; 1994-317007/39.

Dalboge H, Kofo Heldt-Hansen HP;

316 ACCGCGTGCTGGCGGCGGA 334

Conservative

Local Similarity

Query Match

16;

Best Loca Matches

N

Accaccitecrescesces

20

Dp

à

B

AAX29178 standard; DNA; 20

RESULT 115

AAX2917

(first entry)

18-JUN-1999

AAX29178;

Synthetic. Homo sapiens. WO9907844-A2

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Gaps

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                                                                              New antisense compound targeted to a nucleic acid molecule encoding human mitochondrial phosphoenolpyruvate carboxykinase useful for treating a human with a mitochondrial phosphoenolpyruvate carboxykinase-associated
                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                  AAZ95320 to AAZ95359 represent antisense oligonucleotides targeted to a nucleid acid modecule encoding human mitochondrial phosphoenolpyruvate carboxykinase (also known as PEPCK-mitochondrial, PEPCK-M; PCK2 and mtPEPCK), where the oligonucleotide specifically hybridise with and inhibit the expression of human mtPEPCK. The antisense oligonucleotides can be used for inhibiting the expression of mtPEPCK in human cells or tissues in vitro and can also be used for treating an animal, particularly a human suspected of having or being prone to a condition odisease associated with expression of mtPEPCK. They can also be used in diagnostics and as research reagents in sandwich and other assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, mouse, B7-1, B7-2, antisense, PCR primer, inflammation, autoimmune disorder, phosphorothioate backbone, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.4e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human B7-1 antisense oligonucleotide SEQ ID NO: 154.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Karras JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 12; Page 76; 162pp; English.
  Butler MM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                136 CCCGCCTGGCGGTGGAGGC 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 ccadccrddcadrdcaddc 20
                                                                                                                                                                              Claim 3; Col 39; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAY-2000; 2000WO-US014471.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-00326186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF32957 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vickers TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC.
  Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-049991/06.
                                        WPI; 2000-205209/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200074687-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF32957;
Mckay R,
                                                                                                                                          disease.
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The present invention provides sequences of antisense oligonucleotides targeted at the murine and human B7-1 and B7-2 coding and mRNA sequences. The antisense sequences have phosphorothicate backbones and some nucleotides are 2'-methoxyethoxy residues. The sequences can be used in the treatment of inflammatory and autoimmune disorders, including asthma,

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The invention provides Zea mays signal transduction proteins and encoding nucleotide sequences. The nucleic acids are useful for regulating captured sequences. The nucleic acids are useful for regulating captured specially of phytohormones, including ethylene, auxins, cytokinins, and gibberslin, to effect developmental changes in plants and provide control of plant response to environmental stresses. They may also be used as probes or amplification primers in the detection, quantitation or isolation of gene transcripts, for detecting mutations in the gene, for monitoring upregulation of expression or changes in enzyme activity in screening assays of compounds, for detection of any number of allelic variants, or for site-directed mutagenesis in enkaryotic cells. They may further be used for recombinant expression of their encoded polypeptides, as immunogens in the preparation or screening of antibodies, and in sense or antisense suppression of genes in a host cell, tissue or plant. The proteins may be used in assays for enzyme agonists or antisquists, as simunogens or antisques to obtain antibodies specifically immunoreactive with the proteins. The present sequence represents a PCR primer used for amplifying the cDNA encoding a signal transduction protein
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juvenile diabetes mellitus, myasthenia gravis, Graves' disease, rheumatodi arthritis, allograft rejection, inflammatory bowel disease, multiple sclerosis, psoriasis, systemic lupus erythematosus, contact dermatitis, rhinitis, allergies and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New signal transduction nucleic acids and encoded proteins useful for regulating phytohormone expression, including ethylene, auxins, cytokinins and gibberellin, to provide control of plant response to environmental stresses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     phytohormone; ethylene; primer; ss.
                                                                                                                                                                            Gaps
                                                                                                                                                                            ;
                                                                                                                                     Length 20;
                                                                                                                                                                      3; Indels
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                                                                                              Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                 3.3%; Score 14.2; DB 1;
84.2%; Pred. No. 2.4e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zea mays; maize; signal transduction protein;
auxin; cytokinin; gibberellin; immunogen; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Signal transduction cDNA amplifying primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 111; 126pp; English.
                                                                                                                                                                                                                 398 GAAGGTCTTCTACGTGATC 416
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                                                                                                                                                                                                                                                                                                                                                   AAC84282 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                 3.33
Best Local Similarity 84.23
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-031929/04.
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                                                                                                                                                                                                                                                                                                              RESULT 11
AAC84282/
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Synthetic.
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AAD40675/c
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                                  Gaps
                                                                                                                                                                                                                                                                                                         Human, hepsin, antisense compound; antisense therapy; antisense; phosphorothioate backbone; ss.
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Query Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.4e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                              Human hepsin antisense oligonucleotide, ISIS 107131.
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/mod_base= m5c
16. _20
/*tag= c
/mod_base= OTHER
/note= "2'methoxyethyl nucleotides"
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"tag= b" orHER"
"not == "2'methoxyethyl nucleotides"
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mod_base= OTHER
note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                              location/Qualifiers
                                                                139 GCCTGGCGGTGGAGGCCGG 157
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mod_base= m5c
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mod_base= m5c
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mod_base= m5c
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mod_base= m5c
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/mod_base= m5c
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AAD40857 standard; DNA; 20 BP.
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Synthetic.
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                                                                                                                                                  Novel antisense compound targeted to nucleic acids encoding human hepsin, useful for inhibiting the expression of hepsin in human cells or tissues, and for treating humans having a disease associated with human hepsin.
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/note= "2'methoxyethyl nucleotides"
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3.3%; Score 14.2; DB 1;
Best Local Similarity 84.2%; Pred. No. 2.4e+02;
Matches 16; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                      Claim 3; Page 97; 100pp; English.
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/mod_base= m5c
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'mod_base= m5c
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/mod_base= 1
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(ISIS-) ISIS PHARM INC
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modified_base
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                                                    Cowsert LM;
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The invention relates to antisense compounds targetted to a nucleic acid encoding human receptor interacting protein (RIP)2 to inhibit its expression. Antisense compounds are used for treating diseases associated with RIP2 expression. They are also useful in antisense gene therapy. The present sequence is an oligonucleotide targetted to human RIP2 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               num annibense oligonucleotide that targets regions of a nucleic acid encoding human receptor interacting protein (RIP)2, for treating diseases associated with RIP2 expression.
                                        interacting protein; RIP2; antisense; gene therapy;
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/mod_base= m5c
/*tag= c
/mod_base= OTHER
/mod_base= OTHER
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Best Local Similarity 84.2%; Pred. No. 2.46+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                'note= "2-methoxyethyl (2'-MOE) nucleotides"
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/mod_base= OTHER
/not_e= "Phosphorothioate backbone"
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           Human RIP2 antisense oligonucleotide ISIS #104251
                                                                                                                                      location/Qualifiers
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/mod_base= m5c
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                                            Human, receptor inter
phosphorothioate; ss.
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modified_base
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                                                                                           Homo sapiens.
Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an antisense compound 8-30 nucleobases in length targetted to a nucleic acid molecule encoding human hepsin. The antisense compound inhibits the expression of human hepsin. The antisense compound or the pharmaceutical composition of useful for treating animals and human having a disease or condition associated with the expression of hepsin, e.g. inflammation or tumour prophylaxis (e.g. to prevent or delay infection, inflammation or tumour formation) or as research reagents and kits. The method is useful for modulating, specifically inhibiting the expression of hepsin which may be used in research, e.g to distinguish between functions of various members of a biological pathway. The invention is used in gene therapy. The present sequence is human hepsin antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotides that modulate (particularly inhibit) human hepsin, useful for treating a disease or condition associated with the expression of hepsin, e.g. inflammation or tumor growth.
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3.3%; Score 14.2; DB 1; Length 20;
Best Local Similarity 84.2%; Pred. No. 2.4e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
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/mod_base= m5c
16, .20
/*tag= c
/mod_base= OTHER
/mod_base= OTHER
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'mod_base= m5c
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'mod_base= m5c
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AAD45181/C
ID AAD45181 standard; DNA; 20 BP.
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AC AAD45181;
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T 27-DEC-2002 (first entry)
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Detection or determination of a protein, a fused protein, a DNA, a vector, purification of a target protein, a solid carrier, an epitope peptide, a kit for the detection or determination.
                                                           Anti-human type II DNA topoisomerase alpha antibody-related DNA #38.
                                                                                                  Human; type II DNA topoisomerase alpha antibody epitope; ss.
                                                                                                                                                                                                                                                                              26-DEC-2000; 2000JP-00394675.
                                                                                                                                                                                                                                                                                                                       26-DEC-2000; 2000JP-00394675.
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                      29-NOV-2002 (first entry)
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                                                                                                                                                  Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method (M1) for the diagnosis of dentinogenesis imperfecta type II and/or its accompanying deafness comprising determining the dentin sialophosphoprotein (DSPP) gene, its transcript and/or protein of an individual for comparison of their sequences with the normal equences and judging the individual to have higher risk of suffering from the disease then the normal population. Also described are: (1) treating dentinogenesis imperfecta type III and/or its accompanying deafness by administering a safe and effective dose of normal DSPP and/or DSP protein to patients; (2) drug compositions containing safe doses of DSPP and/or DSP protein; and (3) a regent kit for detecting dentinogenesis imperfecta type II and/or its accompanying deafness containing probes for binding to the mutation site. The DSPP gene and protein sequences have auditory activity. The method (M1), dentin sialophosphoprotein (DSPP) gene and DSP protein are useful for diagnosing and treating imperfecta type II and/or its accompanying deafness. The DSPP gene is located to chromosome 4421. The present sequence represent sequence invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                          Human, dentin sialophosphoprotein precursor; dentin sialophosphoprotein;
DSPP; dentinogenesis imperfecta type II; deafness; auditory;
chromosome 4q21; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Diagnosis of dentinogenesis imperfecta type III and its accompanying deafness using dentin sialophosphoprotein gene and encoded products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 3 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SHAN-) SHANGHAI RES CENT BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hu L;
                                                                                                                                                                                                                     Human DSPP PCR primer SEQ ID NO:15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 12; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             353 CTACAGCGACTTCCTCACT 371
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                                                                                         ABQ73550 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-AUG-2001; 2001WO-CN001292.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-SEP-2000; 2000CN-00125042.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Xiao S, Zhao G,
                                                                                                                                                                            03-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                            WO200258722-A1.
                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                        01-AUG-2002
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                                                                                                                                  ABQ73550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kong X,
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                                     The invention relates to a target protein fused with a polypeptide having an amino acid sequence containing an epitope of anti-human type II DNA topolsomerase alpha antibody and the DNA encoding it. The sequences can be used in a method for the detection or the determination of a target protein in which the target protein is detected or determined by using the reactivity between the target protein and the above fused protein as the index, and also in a method for the purification of a target protein in which the above fused protein is contacted to anti-human type II DNA topolsomerase alpha antibody carried on a solid carrier. This sequence represents DNA encoding an anti-human type II DNA topolsomerase alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, K-ras, PCR primer; probe, capture probe, mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
3.3%; Score 14.2; DB 1; Length 20;
Best Local Similarity 84.2%; Pred. No. 2.4e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Capture oligonucleptide Zip ID#944 oligo #9.
Disclosure, Page 33; 38pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26 CGAGGGCTGGGACGAGAT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGAGAGCTGGGACATAGAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABI93857 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                             antibody epitope
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ABS66287 standard; DNA; 20

RESULT 123
ABS66287/c
ID ABS6621
XX
AC ABS6621

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ABS66287

14-APR-2000; 2000US-0197271P. 

Gerry NP, Favis R, Kliman R; (CORR ) CORNELL RES FOUND INC. Barany F, Zirvi M, WPI; 2002-034366/04.

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English.

The present invention describes a method (M1) for designing capture oligonuclectide probes (I) for use on a support to which complementary coligonuclectide probes (I) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful so detecting infectious diseases caused by bacterial infectious agents of Salmonalia, Listeria monocytogenes and Hamophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and happeralius fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epsterian Barr virus and palasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 1 hydroxylase deficiency, Purner Syndrome and obesity defects.

Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, or method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe present or absence of the target mucleotide sequences. Bala2074 to Althour and the parason in the exemplification of the present on the present of sequences used in the exemplification of the present invention

Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;

0; Gaps Query Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.46+02; Matches 16; Conservative 0; Mismatches 3; Indels

25 CCGAGGCTGGGACGAAGA 43 20 ccerecearaceaceaea 2 ઠ

ABZ98645 standard; DNA; 20 BP 17-OCT-2003 (first entry) AB298645; RESULT 125 ABZ98645/ 

Human tryptase a oligonucleotide sequence.

Human, antisense, lung dysfunction; nasal airway dysfunction, antinflammatory steroid, ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive, immunosuppressive, cytostatic; gene therapy, antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure, SEQ ID NO 13887; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonuclectide antisense to the instantion codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of intitiation codon, confirm a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entitificammatory steroid and ubjournone. A composition of the invention has antiinflammatory antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antimilammatory steroid in a subject, for reducing levels of adenosine or lung surfactant in a subject, for reducing levels of adenosine conjung surfactant in a subject, stating bronchoconstriction, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO.

The inventor of the print of the printed specification, but was obtained in electronic format directly from WIPO. 

Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;

Gaps .. 0 Query Match 3.3%; Score 14.2; DB 1; Length 20; Best Local Similarity 84.2%; Pred. No. 2.4e+02; Matches 16; Conservative 0; Mismatches 3; Indels

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RESULT 126 ADE27892

ADE27892 standard; DNA; 20 BP.

ADE27892;

29-JAN-2004 (first entry)

Human B7-1 targeted oligonucleotide SEQ ID 154.

ss; human; B7-1; inflammatory skin disorder; antisense; psoriasis; conteact dermatitis; atopic dermatitis; seborrheic dermatitis; nummular dermatitis; generalised exfoliative dermatitis; eczema; critical costimulatory molecule.

Synthetic

Homo sapiens.

US2003176374-A1. 

18-SEP-2003.

09-MAY-2001; 2001US-00851871.

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The invention relates to a method of treating an inflammatory skin disorder in an individual by topically applying an antisense compound targeted to a nucleic acid molecule encoding a human B7 protein. The invention is for treating an inflammatory skin disorder in individual. The skin disorder is psoriasis, contact dermatitis, atopic dermatitis, seborrheic dermatitis, mummular dermatitis, generalised exfoliative dermatitis or eczema. The invention effectively modulates critical costinulatory molecules such as the B7 protein. The present sequence represents a human B7-1 targeted oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated DNA molecule encoding cholecystokinin receptor protein - are
                                                                                                                                                                               Treating an inflammatory skin disorder such as psoriasis comprises topically applying an antisense compound targeted to the nucleic acid encoding human B7 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cholecystokinin receptor protein; CCK; gastrointestinal receptor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence of nested PCR primer for cholecystokinin (CCK) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                               3.3%; Score 14.2; DB 1; Length 20; 84.2%; Pred. No. 2.4e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                             Example 12; SEQ ID NO 154; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DEPT HEALTH & HUMAN SERVICE
                                                                                                                            Karras JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             398 GAAGGICITCIACGIGAIC 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 GAAGGIGTICTICGTGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92US-00831248.
92US-00861769.
92US-00928033.
92US-00937609.
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            31-DEC-1996; 96US-00777266.
04-JUN-1999; 99US-00326186.
25-MAY-2000; 2000WO-US014471.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ47676 standard; cDNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                                                            Bennett CF, Vickers TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 84.2
nes 16; Conservative
                                                                   (BENN/) BENNETT C F. (VICK/) VICKERS T A. (KARR/) KARRAS J G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NPI; 1993-272886/34
                                                                                                                                                       WPI; 2003-863863/80.
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11-AUG-1992;
02-SEP-1992;
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07-FEB-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-FEB-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ47676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 127
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Mixed oligos primed amplification of CCK cDNA was performed using 2 groups of degenerate primers based on the AA sequence from AAR38890. The sense gp. of primers was 72 fold degenerate (AAQ47672). The anti- gp. of primers was 80 fold degenerate and consisted of AAQ47673. The anti- gp. of primers was 80 fold degenerate and consisted of AAQ47673 & AAQ47674. The product of the PCR was used to generate nondegenerate primers for subsequent PCR. The remaining 3' coding and UTRS was obefor using amplificn. (RACE) of CDNA and anchored PCR. RACE was performed using AAQ47675 for the first round and a nested primer, AAQ47676, for the second round. Anchored PCR used the gene specific primer AAQ47677 and the sequences was cloned using PCR. The sense primer was AAQ47677 and sequences was cloned using PCR. The sense primer was AAQ47679 and the antisense primer was AAQ47680. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid segment comprising one of the 10 - 100 bp sequences given in the specification (sequences of a polymorphic site), or the complement of the segment and a method of analysing a nucleic acid comprising determining the base occupying the polymorphic site of the polymorphic fragment sequences are disclosed in the specification. The information obtained from nucleic acid analysis by the method described is useful in diagnosis or monitoring of diseases like cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid segments containing polymorphic sites, or complements and methods of detecting a nucleic acid - for general use including diagnosis and monitoring of diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
isolate cholecystokinin receptor clones and produce anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; polymorphic site; nucleic acid analysis; diagnosis; monitoring; cancer; inflammation; heart disease; CNS disease.
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                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleotide fragment containing polymorphic site, WI-7038.
                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Indels
                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 6 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                 3.3%; Score 14.2; DB 1;
Local Similarity 84.2%; Pred. No. 2.7e+02;
LE 16; Conservative 0: Mismatchar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Berno A;
  purified to isolate cholecystokinin cholecystokinin receptor antibodies
                                                        Example; Page 38; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              241 GCIGCITCCCGGGCICGGC 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 10; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 GCTGCTGCCAGTGCTCGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV67403 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-US004571
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (APFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chee M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-495419/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Gaps

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21 TGGAGGCAAGGTTCGACTG

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This sequence is an inhibitor of the human AUR2 protein of the invention. The AUR1 and AUR2 proteins can be used to identify specific modulators of, and to generate specific antibodies recognising AUR1 and AUR2. The modulators can be used for treating conditions involving abnormal AUR signal transduction, specifically cancer (of colon, breast, kidney, ovary, bladder, head or neck, also glioma, medullablastoma, kidney, chondrosparcoma and panoreatic tumnurs, particularly of colon (specifically), breast or kidney). The modulators can also be used for studying their effects in animal models of proliferative disease. Probes, based on the coding sequences are used, diagnostically, to detect or quantify AUR mRNA, by hybridistation or polymerase chain reaction (PCR). The DNA, optionally mutated, are useful in gene therapy. Ab are used as diagnostic immunoassay reagents for detecting the proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid encoding human AUR1 and 2 polypeptides, used to identify specific modulators for treating cancer or for diagnosis.
inflammation, heart disease, CNS diseases, and susceptibility to infection by microorganisms. In addition, the nucleic acid segments are useful in manufacturing medication in the treatment of prophylaxis of diseases, and also the use of the DNA segments as pharmaceutical
                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AUR1; AUR2; human; AUR modulator; cancer; glioma; medullablastoma; chondrosarcoma; pancreatic tumour; proliferative disease; diagnosis; therapy; inhibitor; ss.
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                                                                                                                             Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 76.2%; Pred. No. 2.7e+02; Matches 16; Conservative 1; Mismatches 4; Indels
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                                                                                            Sequence 21 BP, 3 A, 4 C, 7 G, 6 T, 0 U, 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 24; Page 120; 153pp; English.
                                                                                                                                                                                                               GCCACCACTCAGAGGAGTCTC 66
                                                                                                                                                                                                                                    21 GCCATCACGCRGAAAAGICTC 1
                                                                                                                                                                                                                                                                                                                                               AAX99728 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-US001283
                                                                                                                                                                                                                                                                                                                                                                                                                           29-SEP-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plowman GD, Mossie K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human AUR2 inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-458699/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SUGE-) SUGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9937788-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                        AAX99728;
                                                                                                                                                                                                               46
                                                                                                                                                                                                                                                                                                            RESULT 12:
AAX99728/
                                                                                                                                                                                                                                                                                                                                                                                      8888888
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The present invention describes human mitogen-activated protein kinase/extracellular response kinase (MAPK/ERK) kinase kinase (MEKK). Pepocifically designated MEKKI, MEKKZ and MEKKI. The MEKK proteins are used to modulate and regulate signal transduction in cells, as well as for regulation of gene transcription in a cell encoding MEKK, where the cell is involved in inflammation, regulation of cellular prolliferation and differentiation, regulation of development, regulation of cell death or regulation of inflammation. They are also used to prepare antibodies. MEKK polymucleorides can be used to produce the protein recombinantly and as a source of probes and primers. The present sequence represents a PCR primer for human MEKK2, which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human MEKK polynucleotides and polypeptides, used for regulating signal transduction in cells.
                                                                                                                            MEKKI; MEKK2; MEKK3; mitogen-activated protein kinase; MAPK; ERK; extracellular regulated kinase; signal transduction; regulation; MAPK, ERK; MEK; MEKK; inflammation; cellular proliferation; differentiation; development; cell death; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
3.3%; Score 14.2; DB 1; Length 21;
Best Local Similarity 84.2%; Pred. No. 2.7e+02;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 5 A; 3 C; 11 G; 2 T; 0 U; 0 Other;
                                                                                                Human MEKK2 PCR primer SEQ ID NO:28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 64; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          289 AGCTGGTGAAGGACCTGAG 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                                                                                                                                   98US-0078153P.
AAZ25089 standard; DNA; 21
                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                    (CADU-) CADUS PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA52302 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-571843/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-SEP-2000
                                                                                                                                                                                                                Synthetic.
Homo sapiens.
                                                                09-DEC-1999
                                                                                                                                                                                                                                                                  WO9947686-A2
                                                                                                                                                                                                                                                                                                                                  15-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                    16-MAR-1998;
04-SEP-1998;
                                                                                                                                                                                                                                                                                                  23-SEP-1999,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Johnson GL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA52302;
                                AAZ25089;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA52302
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Gaps

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3.3%; Score 14.2; DB 1; Length 21; llarity 84.2%; Pred. No. 2.7e+02; Conservative 0; Mismatches 3; Indels

Query Match Best Local Similarity Matches 16; Conserv

148 TGGAGGCCGGCTTCGACTG 166

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Plasminogen, human, kringle 5 domain, endothelial cell proliferation, anglogenesis, antiproliferative, antiarteriosclerotic; cytostatic, antiarflammatory; antialcer; antirheumatic; antiarthritic, antiangiogenic; cancer; tumour; autodamune disease; Escherichia coli; recombinant expression; vector construction; PCR primer; ss.
       Oligonucleotide used to construct UpEt-Ubi vector, SEQ ID NO:31.
                                                                                                                          96US-00643219.
                                                                                                            97US-00851350
                                                                                                                                                                              WPI; 2000-349573/30.
                                                                                                                                                (ABBO ) ABBOTT
                                                                                                             35-MAY-1997;
                                                                                                                          03-MAY-1996;
03-APR-1997;
                                                                                                                                                               Davidson DJ;
                                                                               JS6057122-A.
                                                                                              32-MAY-2000
                                                                 Synthetic.
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Preparation of Kringle five peptide fragment for treating various disorders such as angiogenic, ocular, skin diseases and cancer, involves mixing mammalian plasminogen and elastase followed by incubation and isolation. Example 20; Col 49; 48pp; English

The invention relates to a method of preparing plasminogen kringle 5

peptide fragments. The method comprises mixing mammalian plasminogen and
elastase in the ratio 1:100-1:300, followed by incubating and isolating
the fragment. The kringle 5 peptides are inhibitors of angiogenesis and
endothelial cell proliferation and migration. The peptides are useful for
treating angiogenic diseases, primary and metastatic solid tumours and
carcinomas of various organs such as breast, genital tract, endocrine
glands, skin, tumours of the brain and eyes and solid tumours arising
are also used for the prophylaxis of various autoimmune diseases (e.g.,
rheumatoid arthritis), ocular diseases, skin diseases (e.g.,
blood vessel diseases (e.g. haemangiomas, Osler-Webber Syndrome),
creation of endothelial cells
(e.g., Crohn's disease, atherosclerosis), diseases which have
angiogenesis as a pathologic consequence (e.g., cat scratch disease and
clers). The peptides are also useful as a birth control agent which
inhibits ovulation and establishment of the placenta. Sequences AAA52294AS2304 represent PCR primers used in the construction of Escherichia cells
expression vectors for recombinant expression of various human plasminogen kringle 5 fragments

Sequence 21 BP; 7 A; 6 C; 8 G; 0 T; 0 U; 0 Other;

Gaps ö 3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels Conservative Query Match Best Local Similarity Matches 16; Conserv

CCGCGACGACGCCCAAG 398 21 ceeceaceaceaceaceae 380 m

AAA52303 RESULT

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AAA52303 standard; DNA; 21 (first entry) 18-SEP-2000 AAA52303; KAKKA

human; kringle 5 domain; endothelial cell proliferation; Plasminogen; 

Oligonucleotide used to construct UpEt-Ubi vector, SEQ ID NO:32

angiogenesis, antiproliferative, antiarteriosclerotic, cytostatic, antipsortatic, antilnflammatory, antilnflammatory antilnflammatory antilnflammatory antilnflammatory antilogenic, cancer, tumour, autoimmune disease, Escherichia coli; recombinant expression, vector construction, PCR primer, ss.

Synthetic.

US6057122-A.

02-MAY-2000

97US-00851350 05-MAY-1997; 96US-00643219. 97US-00832087. 03-MAY-1996; 03-APR-1997;

LAB. (ABBO ) ABBOTT

Davidson DJ;

WPI; 2000-349573/30.

Preparation of Kringle five peptide fragment for treating various disorders such as angiogenic, ocular, skin diseases and cancer, involves mixing mammalian plasminogen and elastase followed by incubation and isolation

Example 20; Col 49; 48pp; English.

The invention relates to a method of preparing plasminogen kringle 5
clastase in the ratio 1:100-1:300, followed by incubating and solating
the fragments. The method comprises mixing mammalian plasminogen and
elastase in the ratio 1:100-1:300, followed by incubating and solating
the fragment. The kringle 5 peptides are inhibitors of angiogenesis and
endothelial cell proliferation and migration. The peptides are useful for
treating angiogenic diseases, primary and metastatic solid tumours and
carcinomas of various such as breast, genital tract, endocrine
glands, skin, tumours of the brain and eyes and solid tumours arising
from haematopoietic malignancies such as leukaemias and lymphomas. They
are also used for the prophylaxis of various autoimmune diseases (e.g.,
rheumatoid arthritis), coular diseases, skin diseases (e.g.,
plood vessel diseases (e.g. haemangiomas, Osler-Webber Syndrome),
cliseases caused by axcessive or abnormal stimulation of endothelial cells
(e.g., Crohn's disease, atherosclerosis), diseases which have
angiogenesis as a pathologic consequence (e.g., cat scratch disease and
ulcers). The peptides are also useful as a bitth control agent which
chibits ovulation and establishment of the placents. Sequences AAA52294.
A22304 represent PCR primers used in the construction of Escherichia coli
expression vectors for recombinant expression of various human fragments kringle 5

Sequence 21 BP; 0 A; 8 C; 6 G; 7 T; 0 U; 0 Other;

Gaps ö 3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; tive 0; Mismatches 3; Indels Local Similarity 84.2 les 16; Conservative Query Match Best Loca

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380 CCGCGACGACGCCCAAG 398 19 ceeceaceaceaceaea

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ВЪ AAF29947 standard; DNA; 21 (first entry) 05-APR-2001 AAF29947; RESULT 133
AAF29947/c
XX
AC AAF299
XX
AC AAF299
XX
XX
DE Primer

Primer #5

WO200118250-A2

92US-00831248. 92US-00861769. 92US-00928033. 92US-00937609.

93US-00029170

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The present invention relates to a cholecystokinin (CCK) receptor protein. The CCK receptor-encoding DNA molecule is useful for expressing and purifying CCK receptor protein to sequenceable-grade homogeneity. The CCK receptor proteins or fragments are useful for obtaining antibodies that can recognize CCK-expressing cells. The transformed eukaryotic cell lines are useful for studying the receptor in an environment similar to its native environment, e.g. in the context of studying the electrophysiology or binding properties of the receptor. The transformed prokaryotic or insect cell line is useful for expressing CCK receptor to produce large amounts of the receptor for immunological purposes or for studying protein structure, e.g. crystallography
                                                                                                                                                                                                                                                                                                                                                               New cholecystokinin (CCK) receptor-encoding DNA molecule, useful for producing and purifying human CCK receptor protein to sequenceable-grade
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myceardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 2.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 6 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human gene single nucleotide polymorphism #895.
              Cholecystokinin; CCK receptor; purify; ss.
                                                                                                                                                                                                                                                              (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Col 11; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241 GCTGCTTCCCGGGCTCGGC 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 derderdecendrechedec 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF96134 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                 WPI; 2001-136725/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-JUN-2001
                                                                                                                                                                                            01-APR-1992;
11-AUG-1992;
02-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                              Unidentified
                                                                              US6169173-B1
                                                                                                                                               10-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                 producing an
homogeneity.
                                                                                                              02-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF96134;
                                                                                                                                                                                                                                                                                               Wank SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF96134
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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various obly morphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, concary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                           Mccarthy JJ;
                                                                                                                                                                                                                                              Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
replace(11,C)
/*tag= a /
/tandard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
                                                                                                                                                                                           Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 1 A; 8 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human gene single nucleotide polymorphism #1853.
                                                                                                                                                                                         Ireland JS, Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                               RES.
                                                                                                                                               (WHED ) WHITEHEAD INST BIOMEDICAL (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                               Example; Page 111; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTGGCCCGCCTGGCGGTGG 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTGGCCGACCTGGCCGTGG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
                                                                                  10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                                                      07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF97092 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                           Lander ES, Gargill M,
                                                                                                                                                                                                                         WPI; 2001-226749/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200118250-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JJN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAR-2001
                           15-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF97092;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF97092,
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Gaps

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/\*tag= a /standard name= "single nucleotide polymorphism"

Location/Qualifiers

(first entry)

replace (11, T)

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Mccarthy JJ;

ES,

ander

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various opportantly objerception of the within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                         Mucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New use of irinotecan for preparation of compositions for treating in subject having genome with variant allele comprising cytochrome subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma;
cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide
cytostatic; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.2; DB 1; Length 21;
Pred. No. 2.7e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:1.
                                                                                                         Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 4 A; 3 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                         Bolk S,
                                         (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                         JS,
                                                                                                                                                                                                                                                                                                    Example; Page 192; 242pp; English.
                                                                                                           Ireland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17 GCGGGTGACCGAGGGCTGG 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 GTGGGTGACCCAGGGTGG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JUL-2002; 2002WO-EP008219
16-AUG-2000; 2000US-0225724P
                                                                (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACF62200 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 84.2
hes 16; Conservative
                                                                                                      Gargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ä
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                                                                                                                                                  WPI; 2001-226749/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kerb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003013534-A2.
                                                                                                                                                                                                                                                             atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2003
                                                                                                           Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACF62200;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACF62200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various bolymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPs shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forenaics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                Mccarthy JJ;
                                                                                                                                                                                                                                                                                Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 4 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human gene single nucleotide polymorphism #2100.
                                                                                                                                                                                                o,
                                                                                                                                                                                             Bolk
                                                                                                                             (WHED ) WHITEHEAD INST BIOMEDICAL RES (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                Ireland JS,
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                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 174; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGCGCCACCAAGCTGGTGA 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGTGGCACAAAGCTGATGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF97339 standard; DNA; 21 BP
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26-JUL-2000; 2000US-0220947P.
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                                           10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
    07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 84.2
ses 16; Conservative
                                                                                                                                                                                                Gargill M,
                                                                                                                                                                                                                                      WPI; 2001-226749/23
                                                                                                                                                                                                                                                                                                                                                  atherosclerosis.
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Query Match

Best Loca Matches

21

RESULT 136

AAF97339;

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Gaps

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cancer p450,

15-MAR-2001

Variation

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Disclosure; Page 32; 86pp; English.
\ddot{x} \ddot{x} \ddot{x} \ddot{y} \ddot{y}
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The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (infedipline oxidase), polypeptide 5 (CYPAAS) polymuclectide (II). (I) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate correction or an appropriate derivative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of troatment efficiently dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM36013 represent sequences used in the exemplification of the present invention
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Sequence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other;

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Gaps
                                   ;
0
3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; tive 0; Mismatches 3; Indels
                                     16; Conservative
     Query Match
Best Local Similarity
                                   Matches
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336 GACCAGGGCCGGCTGCTCT 354 GTCCTGGGCCGGCTGCTGT 19 ਨੇ

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ACF62201 standard; DNA; 21
         (first entry)
         08-OCT-2003
     ACF62201;
RESULT 138
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BP.

Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:2.

Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide cytostatic; PCR primer; ss.

5

Synthetic.

WO2003013534-A2.

20-FEB-2003

23-JUL-2002; 2002WO-EP008219.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Heinrich G, Kerb R;

WPI; 2003-268144/26.

New use of irinotecan for preparation of compositions for treating in subject having genome with variant allele comprising cytochrome subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.

Disclosure; Page 32; 86pp; English.

The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or panoreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (infedipine oxidase), polypeptide 5 (CYP3A5) polynucleotide (II). (I) and (II) have

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cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate desage and/or an appropriate derivative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of drug resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the exemplification of the present invention
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Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;

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Gaps
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/ Match 3.3%; Score 14.2; DB 1; Length 21; Local Similarity 84.2%; Pred. No. 2.7e+02; nes 16; Conservative 0; Mismatches 3; Indels
         Query Match
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336 gaccaggccggcrgcrcr 354 greeresecesecreers 21

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ADB20872 standard; DNA; 21 RESULT 139 ADB20872/

BP.

(first entry) 20-NOV-2003

ADB20872;

irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene; MRP1 based cancer related nucleic acid SEQ ID NO:2.

Unidentified.

WO2003013533-A2

20-FEB-2003.

23-JUL-2002; 2002WO-EP008200.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

ď Heinrich G, Kerb WPI; 2003-354397/33.

Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide. 

Disclosure; Page 41; 100pp; English.

The present invention describes a method for the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) golynucleotide (II). (I) has cytostatic activity. (I) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human (preferably African or Asian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.

Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;

RESULT 140

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This invention relates to a novel isolated DNA molecule encoding a cholecystokinin (CCK) receptor protein. The invention also discloses a method for purifying a CCK receptor by solubilising a biological preparation containing CCK receptor in 1% digitionin, applying the solubilised receptor preparation to a cationic exchange resin and purifying the eluate of the resin. The purified eluate is then added to an agarose-bound lectin and applied the eluate to a cibacron blue sepharose column and a CCK receptor protein of sequenceable-grade purity. The CCK receptor protein of the invention may have immunomodulatory activity. The DNA molecule of the invention may have immunomodulatory activity. The DNA molecule of the invention is useful for purifying CCK receptor protein to sequenceable-grade homogeneity. The CCK proteins are useful for neuroendozine modulation of the immune system, and for obtaining antibodies that a recognise CCK-expressing cells. The present sequence represents a RACE PCR primer used to amplify the 3' end of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated DNA molecule encoding a cholecystokinin (CCK) receptor protein, useful for neuroendocrine modulation of the immune system, and for obtaining antibodies that can recognize CCK-expressing cells.
                                                                                                                                                                Rat; ss; PCR; primer; CCK; cholecystokinin receptor; immunomodulator; RACE: rapid amplification of cDNA ends.
                                                                                                                      RACE oligonucleotide #2 used to amplify rat CCK cDNA 3' sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence represents a RACE PCR primer used to amplify the 3' end (Rat cholecystokinin (CCK) receptor cDNA sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; rive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 6 A; 7 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              241 GCTGCTTCCCGGGCTCGGC 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 6; 83pp; English.
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92US-00861769.
92US-00928033.
92US-00937609.
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  ACD26205 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                99US-00443745
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                                                                                    (first entry)
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02-SEP-1992;
10-MAR-1993;
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                                                                                    13-SEP-2003
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                                                                                                                                                                                                                                       Rattus sp.
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                                           ACD26205;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wank SA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method for the use of irinotecan (1) or treating colorectal, cervical, gastric, lung, ovarian or pancreation for cancer, or malignant glioma in subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRR1) polynucleoctide (II). (1) has cytostatic activity. (1) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic ancer, or malignant glioma in a subject, where the subject is a human (preferably African or Asian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                              irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
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44.2%; Pred. No. 2.7e+02;
ve 0; Mismatches 3; Indels
Score 14.2; DB 1; Length 21; Pred. No. 2.7e+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                       MRP1 based cancer related nucleic acid SEQ ID NO:1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        336 GACCAGGCCGGCTGCTCT 354
                                                                                         336 GACCAGGGCCGGCTCT 354
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24-MAY-2002; 2002EP-00011710.
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Best Local Similarity 84.2%;
Matches 16; Conservative
  3.3%;
                                                                                                                      GTCCTGGCCGGCTGCTGT
                                                                                                                                                                                                                                           ADB20871 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                             (first entry)
Query Match
Best Local Similarity 84.2'
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified.
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Kerb R;
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                                                                                                                                                                                                                      Heinrich G,
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ADB96944/c
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     ਨੇ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTAl gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTAL gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant gliona. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGT1A1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGT1A1 gene product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I inhibitor; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member A1.
                                            ss; irinotecan; cancer; UGTIA1; cytostatic; topoisomerase I inhibitor; colocecal cancer; cervical cancer; astric cancer; lung cancer; covarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human UGT1A1 variant allele sequence fragment SEQ ID NO:1.
Human UGT1A1 variant allele sequence fragment SEQ ID NO:2.
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24-MAY-2002; 2002EP-00011710.
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The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTAL gene, and if the patient has one or more variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTALA gene. The invention has cytostatic activity. A composition of the invention acts as a topolsomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant gliona. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                                                   Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
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84.2%; Pred, No. 2.7e+02;
tive 0; Mismatches 3; Indels
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                                                                                                                                 (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                Claim 8; Page 44; 107pp; English.
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24-MAY-2002; 2002EP-00011710.
23-JUL-2002; 2002WO-EP008217.
                                                23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
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Best Local Similarity 84.2
Matches 16; Conservative
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Med Apr 21 12:38:21 2004

The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating coloractal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (WDR1) polymolectide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide. Claim 4; Page 69; 130pp; English WPI; 2003-268145/26. 

Sequence 21 BP; 6 A; 9 C; 6 G; 0 T; 0 U; 0 Other;

ò 3.3%; Score 14.2; DB 1; Length 21; 84.2%; Pred. No. 2.7e+02; ive 0; Mismatches 3; Indels 336 GACCAGGGCCGGCTGCTCT 354 0 ო Query Match
Best Local Similarity 84.2
Matches 16; Conservative ò

21 Gréciedecedecreter g

ADB96943 standard; DNA; 21 BP. ADB96943

04-DEC-2003 (first entry)

Human UGT1A1 variant allele sequence fragment SEQ ID NO:1.

irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; panoreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1; TOP1.

Homo sapiens.

WO2003013537-A2.

20-FEB-2003

23-JJL-2002; 2002WO-EP008218.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Kerb R; Heinrich G, WPI; 2003-268145/26.

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polymucleotide.

Claim 4; Page 69; 130pp; English.

The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal cervical, gastric, lung, ovarion or pancreatic cancer, amalignant glioma in a subject having a genome with a variant allele which

ö comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the Gaps ö Length 21; 3; Indels Sequence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other; Score 14.2; DB 1; Pred. No. 2.7e+02; 0; Mismatches 336 GACCAGGCCGGCTGCTCT 354 Match 3.3%; Local Similarity 84.2%; 16; Conservative invention. Query Match Matches 888888888 8

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ADB92134

ADB92134 standard; DNA; 21

ADB92134;

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(first entry) 04-DEC-2003 Human UGT1A1 variant allele sequence fragment SEQ ID NO:1.

irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.

Homo sapiens.

WO2003013535-A2.

20-FEB-2003.

23-JUL-2002; 2002WO-EP008220.

23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710.

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

Heinrich G, Kerb R;

WPI; 2003-342400/32.

New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.

Disclosure, Page 41; 104pp; English.

The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant gloma in a subject having a genome with a variant allele which comprises a multidrug resistance ! (MDR1) polymuclectide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention. 

Seguence 21 BP; 0 A; 6 C; 9 G; 6 T; 0 U; 0 Other;

Gaps ô Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 2.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels Best Loca Matches

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336 GACCAGGGCGGGCTGCTCT 354 GICCIGGCCGGCIGCIGT 19

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                                                                                                                                                                                                                                                             irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
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Best Local Similarity 84.2%; Pred. No. 2.7e+02;
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ADB92135 standard; DNA; 21 BP
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24-MAY-2002; 2002EP-00011710.
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ID ADC2

AC ADC2

XX ADC2

XX ADC2

XX IB-D

DE Huma

XX DOC Huma

XX MOCC

XX MOC
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The invention relates to an isolated or purified polymucleotide encoding a polypeptide (the wild-type form of which is involved in synaptogenesis) that includes at least one mutation associated with development of eneurological disease and/or a predisposition to development of mental contents that modulate their activity. Also mucleic acid, polypeptide, polypeptide agents that modulate their activity. Also mucleic acid, polypeptide, or polypeptide are used to screen for agents that modulate their activity. Also mucleic acid, polypeptide, or host cells containing the vector, are useful as pharmaceuticals for treating mental and neurological disorders, specifically autism, Asperger syndrome, schizophrenia and attention deficit hyperactivity disorder. The wild-type forms of the nucleic acid and polypeptide can be used similarly. Also detecting mutations in the nucleic acid and polypeptide, can be used to disorders that affect formation of syndroms to dispusse and to disgnose mental disease. This sequence corresponds to a PCR primer used to amplify the human wild type HNL4X (ADC24764) and HNL4Y (ADC24764) genes.

New nucleic acid encoding mutant protein involved in synaptogenesis, useful for treatment and diagnosis of e.g. autism, Asperger syndrome, and schizophrenia.

Example 1; SEQ ID NO 21; 416pp; French.

Leboyer M;

Quach H, Betancur C,

Jamain S,

Bourgeron T, Gillberg C;

WPI; 2003-493399/46.

(INSP ) INSERM INST NAT SANTE & RECH MEDICALE. (INSP ) INST PASTEUR. (ASSI-) ASSISTANCE PUBLIQUE HOPITAUX PARIS.

30-NOV-2001; 2001CA-02364106 02-DEC-2002; 2002WO-FR004134

WO2003045998-A2

05-JUN-2003

Homo sapiens

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84.2%; Pred. No. 2.7e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                       Sequence 21 BP; 3 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                  41 AGATGGCCACCACTCAGAG 59
                                                                                                                                                                                                                           20 AGAAGGCCATCATTCAGAG 2
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Best Local Similarity
"" 16; Conservative
"" 16; Conservative
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RESULT 149 ADE77842/c

ss; gene; atherosclerotic lesion; antiatherosclerotic; cerebroprotective; antianginal, thrombolytic; cardiant; ophthalmological; neuroprotective; nephrotropic; vasotropic; atherosclerosis; stroke; angina; thrombosis; myocardial infarction; ischaemic heart disease; transplantation-induced sclerosis; intermittent claudication; diabetes; peripheral artery disease; congestive heart failure; retinopathy; neuropathy; thrombosis. DNA oligo (SeqID 93) encodes peptide that binds atherosclerotic lesions 멾. ADE77842 standard; DNA; 21 (first entry) 29-JAN-2004 ADE77842; 

Synthetic

nootropic; neuroleptic; tranquillizer; gene therapy; synaptogenesis; mutation; neurological disease; mental disorder; psychiatric illness; autism; Asperger syndrome; schizophrenia; autism; Asperger syndrome; schizophrenia; attention deficit hyperactivity disorder; ds; ss; primer.

WO9618736-A2

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This invention relates to novel isolated peptides that selectively bind to mammalian atherosclerotic lesions and as such can be used to detect to mammalian atherosclerotic lesions and as such can be used to detect the in vivo identification of such peptides by using phage display conditions of such peptides by using phage display the peptides. Diagnosis of pathological conditions of the cound by the peptides. Diagnosis of pathological conditions of the endothelial tissue occurs by administration of a peptide conjugated to reporter molecule or therapeutic agent. As such, these peptides can be described variously as antiatherosclerotic, cerebroprotective, antianginal, thrombolytic, cardiant, ophthalmological, neuroprotective, oppides as useful for treating atherosclerosis, as well as identifying the location and severity of an atherosclerosis, myocardial infarction, ischaemic heart disease, transplantation-induced sclerosis and confinence, it is associated with diabetes, which in turn can lead to peripheral artery disease, congestive heart fallomuclectide sequence, isolated from a combinatorial phage display colliboraty, encodes a peptide that binds to atherosclerotic lesions, the aim
                                                                                                                                                                                                                                                                                                                                       Novel peptide which selectively bind to mammalian atherosclerotic lesions, useful for treating atherosclerosis in a mammal, and for identifying location of atherosclerotic lesion in mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 16; SEQ ID NO 93; 286pp; English
                                                                                                                                                                                                                                         Liu C, Edgington TS, Prescott MF;
                                                                                                                                                        (NOVS ) NOVARTIS AG.
(NOVS ) NOVARTIS PHARMA GMBH.
(SCRI ) SCRIPPS RES INST.
                                                                              09-AUG-2002; 2002WO-EP008942
                                                                                                                   10-AUG-2001; 2001US-0311507P
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WO2003014145-A2.
                                       20-FEB-2003.
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Query Match 3.3%; Score 14.2; DB 1; Length 21; Best Local Similarity 84.2%; Pred. No. 2.7e+02; Matches 16; Conservative 0; Mismatches 3; Indels Sequence 21 BP; 6 A; 4 C; 4 G; 7 T; 0 U; 0 Other; 230 CAAATCGGGAGGCTGCTTC 248 ઠે

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Gaps

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21 CAAATCAGGAGTCTGATTC 3 g

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AAX64556 standard; RNA; 15 BP. 20-JUL-1999 AAX64556; RESULT AAX6455 

150

(first entry)

Human B7-1 hammerhead ribozyme target SEQ ID NO:1188.

Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation; diagnosis; ss.

HIV-1 Group O isolate HAM112 PCR primer env25R.

15-JUL-1999 (first entry)

Homo sapiens

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The present invention describes a novel enzymatic nucleic acid (ENA)

chaving a hammerhead motif (HM) comprising: (1) at least 5 ribose residues

chaving a hammerhead motif (HM) comprising: (1) at least 5 ribose residues

con inhibit a 2'-C-allyl modifications; and (iv) a 3'-end modification. The ENA's

con inhibit collagenase and stromelysin production in the synovial

con inhibit collagenate and stromelysin production in the synovial

con inhibit collagenathritis or rheumatoid arthritis. The ENA's can also

particularly osteoarthritis or rheumatoid arthritis. The ENA's can also

be used to treat antigen presenting cells of a donor to induce tolerance

con a recipient to an alloantism of a donor to induce tolerance

con a recipient to an alloantism of autoimmune disease, and for

confacting alergies and other inflammatory conditions. The ENA's can also

treating allergies and other inflammatory conditions. The ENA's can also

confacting alergies and other inflammatory conditions the ENA's can also

conceptagin without introducing the non-specific effects upon gene

expression which accompany treatment with retinoids and dexamethasone.

The concentration of ribozyme required to affect a therapeutic treatment

conceptic. The present sequence is used in the exemplification of the

conception of the present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid molecules having a hammer-head motif - used for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                        Stinchcomb DT, Jarvis T, Draper K, Pavco P;
Gustofson J, Usman N, Wincott F, Matulic-Adamic J;
Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .<u>.</u>
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3.3%; Score 14; DB 1; Length 15;
Best Local Similarity 64.3%; Pred. No. 1.4e+02;
Matches 9; Conservative 5; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 166; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX56095 standard; DNA; 18 BP.
                                                                                                                     94US-00354920.
94US-00363254.
94US-003908324.
95US-00426124.
95US-00432874.
95US-000951P.
95US-000951P.
95US-000951P.
95US-000951P.
                                                                                      95WO-US015516
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 GGUCTUCCUACGUGA 15
                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                              Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                     13-DBC-1994,
23-DBC-1994,
17-FEB-1995,
20-AAY-1995,
04-MAY-1995,
07-JUL-1995,
07-JUL-1995,
07-JUL-1995,
                                                                                    22-NOV-1995;
                                                    20-JUN-1996.
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WO9909410-A2.

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The present invention describes (A) an isolated HIV-1 Group O env polypeptide. Also described are: (1) an isolated HIV-1 Group O env polypeptide. Also described are: (1) an isolated HIV-1 Group O env polypeptide as in (A) or (1); (2) a polymuclactide (PN) encoding a polypeptide as in (A) or (1); (3) an antigen construct comprising a first HIV-1 Group O env polypeptide; (4) an antigen construct comprising a fusion of at least one HIV-1 Group O env polypeptide with at least one HIV-1 Group O env polypeptide, (4) an antigen construct comprising a fusion of a first HIV-1 Group O env polypeptide with at least one additional HIV-1 polypeptide; (6) an antigen construct on a first HIV-2 env polypeptide; (6) an antigen construct as in (3)-(6), (8) an expression vector comprising a pN as in (7); (9) a host cell transformed by an expression vector as in (8); and (10) an immunosesay kit for the detection of antibodies to HIV-1 comprising an antigen construct as in (3)-(6). The antigen constructs can be used for the detection of anti-HIV-1 antibodies in test samples. They can also be used to publify HIV polypeptides, for therapy and for detection of HIV publication o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated HIV-1 Group O env polypeptides - used for the detection of anti-HIV antibodies and for the production of antibodies for use in detection, purification and therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-1; HIV-2; immobilised capture reagent; capillary action; screening, antibody; assay; env protein; PCR primer; ss.
HIV; human immunodeficiency virus; antigen; detection; antibody; differentiation; Group O; env; immunogen; immunoassay; ss.
                                                                                                                                                                                                                                                                                                                                                                                                        Hackett JR, Yamaguchi J, Golden AM, Brennan CA, Hickman RK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match 3.3%; Score 14; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 2.1e+02; Matches 14; Conservative 1; Mismatches 1; Indels
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Human immunodeficiency virus 1.
                                                                         Synthetic.
Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                                                                                                                                        (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                         17-AUG-1998;
                                                                                                                                                                                                                                                                                                         15-AUG-1997;
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                                                                                                                                                  WO9909179-A2
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The invention relates to a rapid assay for simultaneous detection and differentiation of antibodies to HIV-1 groups O and M, and HIV-2. The method comprises (a) contacting the sample with a strip containing at each of comprises (a) contacting the sample with a strip containing at least one immobilised capture reagent per analyte and on which the sample conditions sufficient to form capture reagent per analyte complexes, and (b) detecting a visible colour change at the capture reagent site on the strip wherein the capture capent site on the strip wherein the capture comprises a polypeptide shown in AAV66981-84; and that for HIV-1 group M comprises a polypeptide shown in AAV66981. The invention is used to screen patisents for antibodies to HIV-1 types O and M, and HIV-2. The invention will be particularly useful in the invention provides a screening method which is faster and requires less equipment than prior art methods. Sequences AAX37195-X37222 represent primers used for determining the env seqeunce of the HIV-1 coup.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; secreted protein; immunestimulant; immunesuppressant; virucide; antibacterial; antifungal; cytostatic; antiinflammatory; dermatological; antidiabetic; antiasthmatic; antiarthritic; antirheumatic; profozoacide; antityproid; immune deficiency; averte combined immunedeficiency; SCID; infection; HIV; hepatitis; malaria; autoimmune disorder; systemic lupus; connective tissue disease; multiple sclerosis; erythematosis; rheumatoid arthritis; autoimmune pulmonary inflammation; asthma; duillain-Barre syndrome; autoimmune thyroiditis; myasthenia gravis; insulin dependent diabetes mellitus; graft-versus-host-disease; autoimmune inflammatory eye disease; allergy; hybridisation; probe; ss.
                                                                                                                                                                                                                                                                                       New rapid assay for antibodies to HIV-1 groups O and M, and HIV-2 - can
be used in field assay, requiring no electricity and less specialised
equipment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
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87.5%; Pred. No. 2.1e+02;
/ative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted protein clone ye90_1 probe SEQ ID NO:201.
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                                                                                                                                                                                                        RK,
                                                                                                                                                                                                          Hickman R
Devare SG;
                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 70; 104pp; English.
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                                                                                                                              97US-00912129
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Matches 14; Conservative
                                                                                                                                                                                                          Vallari AS, Hackett JR,
Golden AM, Brennan CA,
                                                                                                                                                                                                                                                               WPI; 1999-190224/16.
                                                                                                                                                                     (ABBO ) ABBOTT LAB.
                                                                                                                              15-AUG-1997;
                                                                                           07-AUG-1998;
                                                     25-FEB-1999.
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HIV-1 group 0; env; gpl20; gp41; glycoprotein; monoclonal antibody; immunoassay; positive control; affinity purification; therapeutic; antigen; expression construct; PCR primer; ss.

HIV-1 env PCR primer env25R, SEQ ID NO:77.

(revised)
(first entry)

15-SEP-2003 22-MAY-2000

Human immunodeficiency virus 1; group O isolate HAM112.

WO200004383-A2.

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AAA16618 to AAA16697 encode the human secreted proteins given in AAY94898

to AAY94980, isolated from human adult brain, adult thyroid, adult
retina, foetal arcinoma, adult blood, adult heural, foetal kidney, adult
placenta, adult testis, whole embryo, adult cartilage, kidney, foetal
callt bladder, cDNA libraries. The polymicleotides and proteins are
predicted to have biological activities which would make them suitable
for treating, preventing or ameliorating medical conditions in humans and
animals. The polymicleotides can be used as markers for tissues in which
the protein is preferentially expressed, as molecular weight markers on
Southern gels, and as chromosome markers or tags to identify chromosomes
or to map gene positions. The proteins can be used in the treatment of
immunedeficiency (SCID), as well as viral, bacterial, fungal and other
infections. These infections include human immunodeficiency virus (HIV),
the proteins can be used to treat autoimmune disorders such
as connective tissue disease, mycobacteria, leismania spp., malaria and
candidiasis. The proteins can be used to treat autoimmune disorders as connective tissue disease, multiple sclerosis, systemic lupus
as connective tissue disease, multiple sclerosis, systemic lupus
cuillain-Barre syndrome, autoimmune thyroiditis, insulin dependent
diabetes mellitus, myasthenia gravis, graft-versus-host-disease and
autoimmune inflammatory eye disease. The proteins can also be used to
treat allergic conditions, such asthma. AAA16688 to AAA16774 represent
probes for the human secreted proteins from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polynucleotides encoding secreted proteins, which may have e.g. nutritional, chemokine, immune stimulating or suppressing, hematopoiesis regulating, tissue growth, activin/inhibin antiinflammatory or tumor inhibition activity.
                                                                                                                                                                                                                                                                                                                                                                                                    Collins-Racie LA, Evans C;
Steininger RJ, Spaulding V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 627; 641pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                K, Mccoy JM, Lavallie ER,
D, Treacy M, Agostino MJ,
Clark HF, Fechtel K;
                                                                                                                                                                           98US-0096815P.
98US-0105368P.
99US-0115334P.
99US-0119931P.
99US-0120575P.
99US-0130500P.
                                                                                                                   99WO-US018298
                                                                                                                                                                                                                                                                                                                                                             (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-205979/18.
                                    WO200009552-A1.
Homo sapiens,
                                                                                                                   13-AUG-1999;
                                                                            24-FEB-2000.
                                                                                                                                                                                                                                                         12-FEB-1999
                                                                                                                                                                                                                                                                               18-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                          Merberg D,
Wong GG,
                                                                                                                                                                                                                                                                                                                                                                                                    Jacobs K,
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Novel monoclonal antibodies useful as positive control reagent for detecting human immunodeficiency virus infections and diagnosing, evaluating or prognosing viral disease.

sxample 2; Page 37; 148pp; English.

Scheffel JW, Hackett JR, Tyner JD, Hickman RK;

WPI; 2000-171290/15.

99WO-US015469.

09-JUL-1999;

27-JAN-2000.

14-JUL-1998; 98US-00115171

(ABBO ) ABBOTT LAB.

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The invention relates to anti-HIV-1 group O monoclonal antibodies, which may be used as positive control reagents in immunoassays to detect and differentiate HIV-1 infections. The invention also encompasses a monoclonal antibody which binds specifically to an HIV-1 group O antigen, which has no more than 15% cross reactivity to a corresponding antigen selected from HIV-1 group M antigens and HIV-2 antigens; and a method of using a monoclonal antibody as a positive control reagent in an immunoassay for the detection of anti HIV-1 group O antibodies. The monoclonal antibodies are useful as positive control reagents in an immunoassay involve coupling a monoclonal antibodies. Such immunoassays involve coupling a monoclonal antibodies. Such immunoassays involve coupling a monoclonal antibodies. Such cantibodies of the invention would be used to ensure that the reagents or immunoassays involve coupling a monoclonal antibodies. Such antibodies of the invention would be used to ensure that the reagents provided to detect HIV-1 group O antibody were performing properly. The monoclonal antibodies are also generating chimeric antibodies for therapeutic used. Different synthetic, recombinant or purification of specific HIV-1 group O-derived proteins from the propersor of HIV antigens can be used in combination in assay to diagnose, evaluate, or prognosticate HIV disease condition. The monoclonal antibodies conformate, and amplify cDNA encoding the native env protein of HIV-1 group O, isolate HAM12. Sequences AA290187-290187 represent CC o, isolate HAM12. Sequences AA2901807 represent PCP prognosor AA2901801 represent pCP prognosor AA2901801 represent pCP prognosor of indiance of field)
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Best Local Similarity 87.59
warches 14; Conservative
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Gaps

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0; Indels

3.3%; Score 14; DB 1; Length 18; 100.0%; Pred. No. 2.1e+02;

100.08;

Local Similarity 100. nes 14; Conservative

Ouery Match Matches

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0; Mismatches

AAZ90302/c ID AAZ90302 standard; DNA; 18 BP. XX

RESULT 154

rng.res

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Page

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Sequences AAC92738-C92817 represent antiense oligomucleotides targetted to the heterogeneous nuclear ribonucleoprotein A1 (hnRNP A1) gene, which inhibit isse expression. The antiense oligomucleotides were designed to target different regions of the human hnRNP A1 mRNA, and were analysed for their effect on hnRNP A1 mRNA levels by quantitative real-time PCR. Cor their effect on hnRNP A1 mRNA levels by quantitative real-time PCR. Cor their effect on hnRNP A1 mRNA levels by quantitative real-time PCR. Cor their effect on hnRNP A1 mRNA levels by quantition in the stabilisation, transport and processing (including alternative splicing) of newly synthesised mRNAs. It facilitates the annealing of single-stranded nucleic acids, modulates the binding of snRNPs to RNA intron sequences, and shuttles continuously between the nucleus and the cytoplasm acting as a carrier protein for mRNAs. hnRNP correlating with shortened to a carrier protein for mRNAs. hnRNP correlating with shortened biogenesis, with low levels of hnRNP correlating with shortened telomeres. In addition, hnRNP A1 has also been classified as an apoptosis class engine grotein on the basis that it is specifically cleaved into three fragments during antibody-mediated apoptosis. Due to its ability to control splicing events, particularly donor splice site selection, hnRNP and invention are useful for diagnosis, prevention and treatment of the invention are useful for diagnosis, such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antisense compound targeted to human hnRNP A1 which specifically hybridizes with and inhibits the expression of human hnRNP A1, useful for modulating the expression of hnRNP A1 in cells.
                                                                     Human hnRNP A1; heterogeneous nuclear ribonucleoprotein A1; heterogeneous nuclear ribonucleoprotein core protein A1; p40CRS; mRNA processing; transport; stabilisation; alternative splicing; donor splice site selection; alternet biogenesis; oncogenesis; apoptosis-associated protein; cancer; tumour formation; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
                         Human hnRNP Al phosphorothioate antisense oligonucleotide, SEQ ID NO:57.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-090484/10.
                                                                                                                                                                                                                                                                                                                                                                                                                   27-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-OCT-1999;
                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3056, 3157, 327, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                           Human biallelic marker downstream amplification primer SEQ ID NO:8409.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                        Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; phybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 5 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 8; Page 2023; 2745pp; English
                                                                        AAZ74053 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-0082614P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-IB000822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0109732P
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                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09954500-A2
                                                                                                                                                                            10-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-OCT-1999
                                                                                                                          AAZ74053;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       205
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AAC92785/c
ID AAC9278
XX
AC AAC9278
XX
DC AAC9278
                RESULT 155
AAZ74053/C
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Gaps

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pulmonary embolism; paternity test; ds

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Page

The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism apulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification Probe; quantification; human; GTP binding protein; G protein; alpha subunit; specific mRNA; detection; hybridisation; diagnosis; pathophysiology; disease state; hereditary; cancer; infectious; osteodystrophy; pituliary tumour; acromegaly; melanoma cells; diabetes; PCR; polymerase chain reaction; ss. Mccarthy JJ; Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis. /standard\_name= "single nucleotide polymorphism" ö 3.3%; Score 14; DB 1; Length 21; 100.0%; Pred. No. 3e+02; ative 0; Mismatches 0; Indels S, Daley GQ, Mouse D MUSJUNDA, MUSJUNDR/B-1258 jun-B specific probe. Sequence 21 BP; 5 A; 4 C; 6 G; 6 T; 0 U; 0 Other; Lander ES, Gargill M, Ireland JS, Bolk AA047598 standard; cDNA to mRNA; 17 BP. (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC. Location/Qualifiers replace(11,T) Example, Page 218; 242pp; English. 10-SEP-1999; 99US-0153357P. 26-JUL-2000; 2000US-0220947P. 16-AUG-2000; 2000US-025724P. 07-SEP-2000; 2000WO-US024503 93 ATCACCACGTCTGA 106 (revised)
(first entry) 19 ATCACCACGTCTGA 6 Query Match
Best Local Similarity 100..
Best Local 14; Conservative WPI; 2001-226749/23. WO200118250-A2 Homo sapiens 25-MAR-2003 26-JAN-1994 15-MAR-2001 Key Variation AAQ47598; RESULT 159 AAQ47598/c g ö Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; Gaps

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The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bolk S, Daley GQ, Mccarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                                                   Sequence 21 BP; 7 A; 4 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                              (WHED ) WHITEHEAD INST BIOMEDICAL RES. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gargill M, Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 184; 242pp; English.
                                                                                                                                                                                                                                                                                                                            10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-025724P.
                                                                                                                                                                                                                                                                                             07-SEP-2000; 2000WO-US024503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-226749/23.
                                                                                                                                                                                                                           WO200118250-A2
                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                            15-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lander ES,
                                                                                                                                                       Variation
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Gaps

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Query Match 3.3%; Score 14; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 3e+02; Matches 14; Conservative 0; Mismatches 0; Indels

ð 원 Human gene single nucleotide polymorphism #2509.

06-JUN-2001 (first entry)

AAF97748/ XX XX AC AAF9 XX XX DT 06-J DT 06-J XX XX XX XX Huma XX KW Puma KW Puma

AAF97748;

748/c AAF97748 standard; DNA; 21 BP.

RESULT 158

WO9954496-A2

28-OCT-1999

99WO-EP002614 19-APR-1999; 98EP-00870088 20-APR-1998;

(INNO-) INNOGENETICS NV

WPI; 1999-634008/54.

Claim 16; Page 19; 62pp; English.

Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA.

Example 9; Page 71; 177pp; English.

Akitaya T, Cooper A, Mitsuhashi M;

WPI; 1993-258695/32.

(HITB ) HITACHI CHEM CO LID. (HITB ) HITACHI CHEM RES CENT INC.

92US-00827208. 92US-00857059. 92US-00974409.

29-JAN-1992; 24-MAR-1992; 12-NOV-1992;

93WO-US000977

29-JAN-1993; 05-AUG-1993

WO9315221-A1

Synthetic.

The invention provides polymucleotides corresponding to exon 2 and exon 3 of human leukocyte antigen (HLA) alleles HLA-B\*3913, HLA-B\*1406 and HLA-B\*510 and exon 2 of HLA alleles HLA-D\*B1\*0620 and HLA-B\*510 and exon 2 of HLA alleles HLA-D\*B1\*0620, HLA-D\*B1\*04 and HLA-B\*510 and exon 2 of HLA alleles HLA-D\*B1\*0620, HLA-D\*B1\*06 and HLA-B\*10. The polymucleotides are useful for typing the above HLA alleles in a sample, especially by a method that comprises (a) amplifying all/part of the relevant sequence using at least one primer pair; and (b) hybridizing to target regions comprising one or more polymorphic nucleotides of the sequence, to determine the absence or presence of the angle in the sample. Diagnostic kits for (a) typing the alleles comprising at least one preferred primer and/or at least one preferred polymucleotides, comprising an antiserum or ligand (e.g. antibody) binding specifically to the protein fragment are provided. The polymucleotides also enable the isolation of the complete respective genes from a human genomic library

Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

The sequences given in AAQ47594-603 show regions of homology between jun sequences and the jun-B specific probe B-1258 which may be of use as jun-B specific probes B-1258 which may be of use as jun-B specific probes. They were used in the method of the invention for the detection and quantification of markha in a sample without the need to purify the mRNA from cells. The claimed method comprises identifying a polymucleotide sequence unique to the mRNA, and immobilising an oligomer complementary to this sequence to an insoluble support. The sample is components are washed from the support such that the unique sequence will hybridise to the bound oligomer and bound RNA is labelled in such components are washed from the support and bound label is then camount of mRNA on the support. The amount of bound label is then amount of mRNA on the support. The amount of bound label is then determined. This method can be used for the reliable, rapid, simultaneous quantification of multiple varieties of mRNA. It may be used for diagnosing and recognition of pathophysiology of various disease states, eg. hereditary diseases, cancer, and infectious diseases. G proteins are thought to be involved in causing various diseases gates. A genetic deficiency of Gs protein is the molecular basis of hereditary costeodystrophy. Pituitary tumours in acromegalic patients have been shown conform multiple washed sproteins are also involved in invasive and metastatic melanoma cells, and diabetes. See also AAQ47381-666.

(Updated on 25-MAR-2003 to correct PN field.)

Local Similarity 88.2 ses 15; Conservative Matches

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AGGACCTGAGCTCCTGG 17

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Gaps ö

Ouery Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels

141 CTGGCGGTGGAGGCCGG 157

17 CTGGCGGTGGACGCCAG 1

Вb

Seguence 17 BP; 2 A; 8 C; 5 G; 2 T; 0 U; 0 Other;

Hammerhead ribozyme substrate #3478.

Human leukocyte antigen; HIA; allele; HIA-B\*3913; HIA-B\*1406; human; HIA-B\*51; HIA-DRB1\*0820; HIA-DRB1\*04; HIA-DRB4\*01; allele typing; exon; major histocompatibility complex; MHC; probe; 98.

Synthetic

Probe for typing HLA allele B\*1406.

SXXXXXXXXXXXXXX

11-FEB-2000 (first entry)

AAZ39286;

AAZ39286 standard; DNA; 17 BP.

RESULT 160

Homo sapiens 

Rossau R; Mersch G, De Canck I, New polynucleotides for human leukocyte antigen, HLA, allele fragments, useful for typing HLA alleles.

3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.1e+02; Artive 0; Mismatches 2; Indels Query Match

298 AGGACCTGAGCCCCGGG 314

Ribozyme; erythropoietin; granulocyte colony stimulating factor;

rng.res

Mcswiggen J;

Pavco P,

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Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                        present invention relates to enzymatic and antisense nucleic acid
                                                                                                                                                                                                                                                                                Claim 54; Page 136; 164pp; English.
                                                                WPI; 2000-647423/62.
ij
Blatt
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molecules that act as inhibitors of the expression of repressor genes encoding the TR2 orphan receptor, EAR3/COUP-TR-1, the GATA transcription factor gene, IRF-2 and/or the CAATI Displacement Protein (CDP) Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha Score 13.8; DB 1; Length 17; Pred. No. 2.1e+02; 0; Mismatches 2; Indels Sequence 17 BP; 2 A; 10 C; 3 G; 2 T; 0 U; 0 Other; ; 0 3.2%; Query Match
Best Local Similarity 88.2.
These 15; Conservative

17 GGGGGACCGAGGGCTTG 1 19 GGGTGACCGAGGGCTGG δ g

ABK00841 standard; RNA; 17 BP (first entry) Human NOGO Inozyme #111. 12-MAR-2002 ABK00841; RESULT 1 ABK00841

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; carebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropachy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. 

WO200159103-A2. sapiens 16-AUG-2001. Synthetic.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. 09-FEB-2001; 2001WO-US004273

RIBOZYME PHARM INC. BLATT L. MCSWIGGEN J. CHOWRIRA B M. (RIBO-) I (BLAT/) I (MCSW/) N CHOM/)

Chowrira BM; Mcswiggen J, 'n, Blatt

Expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGCD). The creation and processing and with a NRW motify processing an Indoxyme or a nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acids desaving an RNA molecule possessing an NCH motify, a d-cleaver (cleaving RNA with a NRW motify) processing an NCH motify and cleaving RNA with a NRW motify processing an NCH motify and cleaving RNA with a NGN triplet), a zinzyme (cleaving RNA with a STAT motify). The CD20-targetting nucleic acid is used to cleave RNA of cofficion in the presence of a divalent cation that is preferably Mg^2+.

C Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more theoremia, HIV (human immunodyform) low-grade or follicular NHL, lymphocytic lymphoma, leukaemia, HIV (human immunodeficiency virus) associated WHL, mantle-cell lymphoma (NCL), immunocytona (MMC), small B-cell lymphocytic lymphoma, immunocytona (MMC), small B-cell lymphocytic lymphoma (NCL), immunocytona (MMC), small B-cell lymphocytic lymphoma (NCC), immunocytona (MMC), small B-cell lymphocytic lymphoma (NCC), immunocytona (MMC), small B-cell lymphocytic lymphocytic catid may be contacted with a cell to reduce NOGO gene in the cargetting nucleic acid may be contacted with a cell to reduce NOGO gene in the NOGO. The treatment may further comprise the use of one or more contact central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzahemer's disease, dementia, multiple sclerosis (MS), charapies. In particular, the NOGO-targetting nucleic acid may be used to cleave which respond to the modular disease, central disease, dementia, multiputed neuropathy, and/or other neurodegenerative of steases which re Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and sequence is an inozyme of the invention Claim 88; Page 79; 200pp; English. central nervous system injury. 

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Gaps . 0

ô Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 1 A; 9 C; 7.G; 0 T; 0 U; 0 Other;

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302 CCTGAGCCCCGGGGACC 318 1 cceccecceceses

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ABN05998 standard; DNA; 17 BP. (first entry) 29-MAY-2002 RESULT 163 ABN05998/C 

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5990.

WO200192524-A2 Homo sapiens

06-DEC-2001

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The present invention describes a human genome-derived myosin-like correction; (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 concleic acids can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify concleic acids can be used as probes to detect, characterise and quantify convide initial substrates for the recombinant engineering of hGDMLP-1 proteins are proteins. The hGDMLP-1 proteins or polypeptides may be expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific blomolecule and/or amount specifically of hGDMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as therappeauting specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concurration and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concurration and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed capture. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO capture.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
             Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                              04-02T-2000; 2000GB-00024263.
30-JAN-2001; 2001MO-US000661.
30-JAN-2001; 2001MO-US000662.
30-JAN-2001; 2001MO-US000663.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
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27-SEP-2000; 2000US-0236359P
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                                                                                                                                                                                  WO200192524-A2
                                                                                                                                  Homo sapiens.
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The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed of the sequence data for this patent did not form part of the printed of the printed in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shannon ME;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 6 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rank DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hanzel DK,
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
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30-JAN-2001; 2001WO-US00670.
05-FEB-2001; 2001US-0266860P.
                                                          2000US-023456P.
2000US-0234687P.
2000US-0235359P.
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2001WO-US000661
2001WO-US000662
        25-MAY-2001; 2001WO-US016981
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                                                                                                                                     04-OCT-2000;
30-JAN-2001;
30-JAN-2001;
                                                          26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
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3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.18+02; ive 0; Mismatches 2; Indels

353 CTACAGCGACTTCCTCA 369

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Local Similarity les 15; Conserv

Matches

Query Match

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17 Chacargaachrechea 1

ABN07570 standard; DNA; 17 BP

RESULT 166

29-MAY-2002 (first entry)

ABN07570;

Sequence 17 BP; 5 A; 2 C; 6 G; 4 T; 0 U; 0 Other;

at ftp.wipo.int/pub/published\_pct\_sequence

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The present sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence in the exemplification of the present invention. N.B. hGDMLP-1 sequence data for this patent did not form part of the princed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polypeptide, for raising antibodies that recognize hGDWLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDWLP-1.
                                                                                                                                                                hGDMLP-1; heart;
heart disease;
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                                                                                                                         Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5989.
                                                                                                                                                                Human, genome-derived myosin-like protein 1; GDMLP-1; Imuscle, myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
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30-JAN-2001; 2001WO-US0006663.
30-JAN-2001; 2001WO-US0006663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US0006666.
30-JAN-2001; 2001WO-US000666.
ABN05997 standard; DNA; 17 BP
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2001WO-US000669.
2001WO-US000670.
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2000US-0236359P.
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2001WO-US000661.
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                                                                                 29-MAY-2002 (first entry)
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30-JAN-2001;
05-FEB-2001;
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27-SEP-2000;
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                          Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                             Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7562.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chen W,
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30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WG-US000670.
05-FEB-2001; 2001US-0266860P.
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30-JAN-2001; 2001MC-US000661.
30-JAN-2001; 2001MC-US000662.
30-JAN-2001; 2001MC-US000663.
30-JAN-2001; 2001MC-US000664.
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27-SEP-2000;
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDMLP-1.
and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as productive supplement in patients having specific deficiancy in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosting a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fitp.wipo.int/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease;
skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                                   Query Match

3.2%; Score 13.8; DB 1; Length 17;

Best Local Similarity 88.2%; Pred. No. 2.1e+02;

Matches 15; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                   Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other;
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21-SEP-2000; 2000US-0234687P.
21-SEP-2000; 2000US-0234585P.
04-0CT-2000; 2000US-0024563.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
                                                                                                                                                                                                                                                                                                                                           387 GACGCCCCAAGAAGGT 403
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ABN05999 standard; DNA; 17 BP.
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ABN05999/c
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Disclosure; SEQ ID NO 5991; 214pp; English

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to hGDMLP-1 covide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically of hGDMLP-1 proteins, as specifically of hGDMLP-1 candor amount specifically of hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as the replacement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration disorders associated with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders, hGDMLP-1 in particular heart cand skeletal muscle disorders, hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part to the printed specification, but was obtained in electronic format directly from MIPO cat figure probablished_pct_sequence
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3.2%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.1e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 6 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-WAY-2001; 2001US-00864761.
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ABV79108
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Wed Apr 21 12:35:21 2004
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The present invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two. One of the single base pair changes introduces a premature stop codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The important in regulating male germ cell development, and the HTPL gene was mapped to human chromosome 10p12.1. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of such disorder associated with decreased expression or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and fortal liver, bone marrow, brain, kidney, lung, placenta, prosette, skeletal muscle or colon function. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potenial therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL. Example 2; Page 110; 718pp; English. 

Sequence 17 BP; 1 A; 8 C; 6 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match 3.2%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.1e+02; Matches 15; Conservative 0; Mismatches 2; Indels

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ABV91035 standard; DNA; 17 BP ABV91035

ABV91035;

23-DEC-2002 (first entry)

Human POSHL1 scanning oligonucleotide SEQ ID NO 1748.

Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss. 

Homo sapiens

EP1239051-A2.

11-SEP-2002.

28-JAN-2002; 2002EP-00001165

30-JAN-2001, 2001WO-US000663. 30-JAN-2001, 2001WO-US000664. 30-JAN-2001, 2001WO-US000665. 30-JAN-2001, 2001WO-US000665. 30-JAN-2001, 2001WO-US000667. 30-JAN-2001, 2001WO-US000669. 30-JAN-2001, 2001WO-US000669. 30-JAN-2001, 2001WO-US000669. 23-MAY-2001, 2001WO-US000670.

(AEOM-) AEOMICA INC.

Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL -1, useful for treating disorders associated with decreased expression or activity of human POSHL1.

The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB83999), a sequence having 65% sequence identity to (S1), (S1), (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequence comprising at least 8 contiguous amino acids. Human POSHL 11s a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GrPsess as well as communities of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) conced by altered expression of human POSHL1 including diagnosing and treating cancer, they useful in the development of vaccines and (II) is treating cancer, they useful in the development of vaccines and (II) is useful for measuring and for surveying gene expression and creating treansgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the prime by the Buropean Patent Office 

Gaps . 0 Query Match
Best Local Similarity 88.2.
These 15; Conservative

ઠ 셤 RESULT 170 ABL31539

Human, human leukocyte antigen, HLA, genotype; polymorphism, immunogenetic, transplantation, genetic disease; ss.

06-DEC-2001.

01-JUN-2001; 2001WO-JP004662.

01-JUN-2000; 2000JP-00164798.

Moriya S, Nishida M; Matsumura Y, Inoko H, Kagiya T, Ichihara T, THE STANDARD STANDARD

Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.

WPI; 2002-684061/74

Example 2; SEQ ID NO 1748; 60pp + Sequence Listing; English.

Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.1e+02; vative 0; Mismatches 2; Indels

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ABL31539 standard; DNA; 17 BP.

21-MAR-2002 (first entry) ABL31539;

Human HLA genotyping oligonucleotide SEQ ID NO 1028.

Homo sapiens

WO200192572-A1.

(NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

WPI; 2002-122074/16.

The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonuclocides (AB10312-AB131809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as priners for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other; xxxyyyyyyyxxx

Claim 10; Page 288; 345pp; Japanese.

3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.1e+02; tive 0; Mismatches 2; Indels 298 AGGACCTGAGCCCCGGG 314 Local Similarity 88.2 es 15; Conservative Query Match

1 Accacinascricación 17

RESULT 171 ABL31778

Human HLA genotyping oligonucleotide SEQ ID NO 1267. ABL31778 standard; DNA; 17 BP (first entry) 21-MAR-2002 ABL31778; 

Human; human leukccyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Homo sapiens.

WO200192572-A1.

06-DEC-2001,

01-JUN-2001; 2001WO-JP004662.

01-JUN-2000; 2000JP-00164798.

(NISN ) NISSHINBO IND INC.

(SYST-) SYSTEM RES INC.

Ichihara T, Matsumura Y, Moriya S, Nishida WPI; 2002-122074/16. Kagiya T, Inoko H,

Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when individuals e.g. by determin transplanting between them.

Claim 10; Page 333; 345pp; Japanese.

The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of clasved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunopenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver,

pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals Sequence 17 BP; 3 A; 5 C; 6 G; 3 T; 0 U; 0 Other; SXSS

3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.1e+02; ive 0; Mismatches 2; Indels Query Match Best Local Similarity

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Gaps

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298 AGGACCTGAGCCCCGGG 314 1 AGGACCTGAGCTCCTGG 17 à

15, Conservative

Matches

7771/c ACA07771 standard; RNA; 17 BP. RESULT 172 ACA0777

ACA0771;

03-JUN-2003 (first entry)

Gaps

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2; Indels

NFKB sub-unit modulating zinzyme substrate #170.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; dc-claaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; besophageal cancer; stomach cancer; close cancer; pencrettic cancer; cervical cancer; necan neck; stomach cancer; ovarian cancer; metanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphands; docetaxel; cisplatin; methotrexate; cyclophosphands; datoremthy; luncouracil carboblatin; datrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; achaemia; transplant/graft respection; reperfusion injury; glomerulonesphritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 18-MAY-1994; 15-AUG-1994; 23-DEC-1996; 07-DEC-1992; 

(STIN/) STINCHCOMB D T. <del>ر</del> DRAPER K G. (MCSW/) I Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 40; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NPKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^22+. The enzymatic and

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prostate, colorectal brain, oesophageal, stomach, bladder, pancreatic, prostate, colorectal brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paciltaxel, docetaxel, cipplatin, methotrexate, gemitabine or radiation therapy. The enzymatic and antisense nucleic coid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomeruhonephritis, september alway inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule

Sequence 17 BP; 2 A; 7 C; 5 G; 0 T; 3 U; 0 Other;

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Gaps
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88.2%; Pred. No. 2.1e+02;
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Best Local Similarity 85.2.
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BD. ADA99411 standard; DNA; 17 ADA99411; 173 ADA99411 

Human MDZ3 scanning oligonuclectide SEQ ID 400.

(first entry)

20-NOV-2003

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ1; MDZ1; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

26-MAR-2001; 2001US-00817879. 08-UUN-2001; 2001US-00877478. 08-UUN-2001; 2001US-0296876P. 24-OCT-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P.

RIBOZYME PHARM INC.

(RIBO-)

BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORKISSEY D.
PAVCO P.
LEE P.
DRAPER K.

26-MAR-2002; 2002WO-US009187

17-0CT-2002.

Hepatitis C virus. WO200281494-A1.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

Gu Y, Nguyen C; Shannon M,

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 400; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD212, MD221, MD212 encoded at chromosome 7q22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15p11.2 and MD212 is encoded at chromosome 15p26.1, The MD23, MD24, MD27, and MD212 sequences are useful in therspy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23,

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MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus; HBV; RNA stability; RNA expression, RNA synchesis; antisense; enzymatic nucleic acid, hammerhead ribozyme; inozyme; inozyme; amberzyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degemerative; disease state; HBV infection; HCV infection; dirrhosis; livar failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                 3.2%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.1e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 3 A; 6 C; 2 G; 6 T; 0 U; 0 Other;
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Best Local Similarity 88.2*
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 174
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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection. Ď Morrissey Мсвиіддеп J, Claim 1; Page 298; 387pp; English Macejak D, Roberts E; WPI; 2003-229207/22 (BLAT) BLATT L.

(MACE) MACEJAK D.

(MCSW) MACEJAK D.

(MCSW) MCSWIGGEN J.

(MORR) MORRISSEY D.

(PAVC) PAVCO P.

(LEEP) LEE P.

(LEEP) RAPER K.

(ROBE) RAPER K. Blatt L, N Draper K,

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HTV) or the synthesis, expression and/or stability of Hepatitis C virus (HTV) or the patitis B virus (HTV) RNA. The nucleic acid and such as hammerhead ribozymes, DNAzymes, and enzymatic nucleic acid such as hammerhead ribozymes. DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. DNAzymes, or are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonuclectides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds that methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence substrate for one of the HCV invention are useful for the treatment of acid season states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular incoming the present sequences disclosed in the present
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3.2%; Score 13.8; DB 1; Length 17; 82.4%; Pred. No. 2.1e+02; ive 1; Mismatches 2; Indels Sequence 17 BP; 5 A; 3 C; 8 G; 0 T; 1 U; 0 Other; 17 76 AGGGCCGCGCAGTGGAC 92 1 AGGCCAGAGCAGUGGAC 14; Conservative Sest Local Similarity Matches ò 셤

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Gaps ; 0

AAZ39244

BP.

Probe for typing HLA allele B\*3913. AAZ39244 standard; DNA; 18 (first entry) 11-FEB-2000 AAZ39244; 

Human leukocyte antigen; HIA; allele; HIA-B\*3913; HIA-B\*1406; human; HIA-B\*51; HIA-DRB1\*0820; HIA-DRB1\*04; HIA-DRB4\*01; allele typing; exon; major histocompatibility complex; MHC; probe; ss.

Homo sapiens. Synthetic.

WO9954496-A2

28-OCT-1999.

99WO-EP002614. 19-APR-1999;

(INNO-) INNOGENETICS NV.

98EP-00870088.

20-APR-1998;

De Canck I, Mersch G,

WPI; 1999-634008/54.

New polynucleotides for human leukocyte antigen, HLA, allele fragments, useful for typing HLA alleles.

Claim 16; Page 18; 62pp; English.

The invention provides polynucleotides corresponding to exon 2 and exon 3 of human leukocyte antigen (HLA) alleles HLA-B\*3913, HLA-B\*1406 and HLA-B\*51 and exon 2 of HLA alleles HLA-DRB1\*0820, HLA-DRB1\*04 and HLA-DRB4\*01. The polynucleotides are useful for typing the above HLA alleles in a sample, especially by a method that comprises (a) amplifying all/part of the relevant sequence using at least one primer pair; and (b)

This oligonucleotide represents a soybean retroelement primer binding site (version 2). The invention provides molecular tools in the form of retroelements and retroelement-containing vectors, cells and plants.

Methods are provided for introducing the retroelements into cells, especially when the retroelement carries at least 1 agronomically. Consideratistic. In a preferred method, a helper cell line which expresses gag, pol and env sequences is used to enable transfer of which expresses gag, pol and env sequences is used to enable transfer of a secondary construct which carries an agronomically significant and integration. Claimed isolated nucleic acid molecules comprise a nucleic acid sequence selected from a retroelement primer binding site, nucleic acid sequence selected from a retroelement primer binding site, envelope, gag, integrates, reverse transcriptase, protease or RNAse-H sequence (see AAZ5254-661). Also provided are plant retroviral particles that are used to transfer the nucleic acids into plant cells ö hybridizing the amplified product to a set of probes specifically hybridizing to target regions comprising one or more polymorphic nucleotides of the sequence, to determine the absence or presence of the allele in the sample. Diagnostic kits for (a) typing the alleles comprising at least one preferred primer and/or at least one preferred probe and (b) for detecting the protein fragment encoded by the polymucleotides, comprising an antiserum or ligand (e.g. antibody) binding specifically to the protein fragment are provided. The polymucleotides also enable the isolation of the complete respective Gaps New nucleic acid molecules for imparting agronomically significant characters to plants, especially soybean. 0 3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; ive 0; Mismatches 2; Indels Retroelement; retrovirus; transgenic plant; gene transfer; Plant retroelement primer binding site version 2. Sequence 18 BP; 3 A; 5 C; 7 G; 3 T; 0 U; 0 Other; Claim 1(a); Page 72; 118pp; English. genes from a human genomic library primer binding site; soybean; ss. 298 AGGACCTGAGCCCCGGG 314 38 98US-0087125P. 99WO-US011858 2 Accaccrcaccrccrcc AAZ35254 standard; DNA; 18 Query Match
Best Local Similarity 88.2'
Matches 15; Conservative Wright DA, Voytas DF; WPI; 2000-105586/09. (WRIG/) WRIGHT (VOYT/) VOYTAS 28-MAY-1999; WO9960842-A2 29-MAY-1998; 28-MAY-1999; 02-DEC-1999. 27-MAR-2000 Glycine max. AAZ35254; 8888888888888 ò 셤

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Human, mouse, Zmax1, HBM, high bone mass gene, lipid regulation, stroke, lipid-associated condition, arteriosclerosis, cardiovascular disease, ss, osteoporosis, atherosclerosis, diabetic arherosclerosis, plaque build-up, neurovascular condition, wound healing; gene therapy, PCR primer, probe, bone development disorder, antiarteriosclerotic, cardiovascular,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method for identifying a molecule involved in inhibits binding of a molecule to high bone mass (HBM) or its wild type gene, Zmaxi. Compounds identified by the method are useful for treating, diagnosing, preventing or screening for normal and abnormal lipid-associated conditions, including arteriosclerosis, cardiovascular disease, stroke, and osteoporosis. The compounds may also be used in the treatment or prevention of diabetic atherosclerosis, neurovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conditions caused by plaque build-up, poor circulation due to plaque build-up and associated poor wound healing. The methods may be used in gene therapy, pharmaceutical development, and diagnostic assays for bone development disorders Molecules identified by comparison of zmaxl and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HBM systems can be used as surrogate markers in pharmaceutical development, in diagnosis of human or animal bone disease, and in the treatment of bone disease. Sequences ABK22776-ABK23411 represent cDNA molecules encoding human Zmax1 and HBM, and PCR primers, probes, linkers and adapters of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying molecules involved in lipid regulation, useful for diagnosing, treating or preventing e.g., arteriosclerosis, compidentifying a molecule that binds to high bone mass gene or its corresponding wild type gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; Volume 18; Nismatches 2; Indels
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                                                                                                                         CDNA reverse PCR primer #226.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 41; 409pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                   osteopathic; cerebroprotective
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                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-097784/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200192891-A2
                                                            09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                         Human Zmax1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-DEC-2001.
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      ABK23290;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; high bone mass; HBM gene; Zmaxl gene; chromosome 11; 11q13.3; sequence tagged site; STS; osteoporosis; osteopathic; gene therapy; antisense therapy; vaccine; bone disorder; Paget's disease; adapter; solerostosis; osteomalacia; fibrous dysplasia; PCR primer; linker; ss.
                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New high bone mass (HBM) and Zmaxl genes and proteins useful for modulating bone mass for the treatment of e.g. osteoporosis.
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                                                         3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zmax1 gene region physical map preparation STS marker #452.
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Sequence 18 BP; 1 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Johnson ML
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                                                                                                                                                                                380 CCGCGACGACGCCCA 396
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                                                                                      15; Conservative
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Synthetic.

Homo

ABA82493;

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Query Match Best Local

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Matches

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Gaps

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(first entry)

19-DEC-2002

197

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RESULT 178

ABK23290 ID ABK2 XX

Query Match

Best Loca Matches

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Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; tranquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropathic pain; theime a disease; glaucoma; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis; memory impairment; renal disease; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy, neuropathic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rossomando
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 8; 138pp; English.
Neublastin DNA related PCR primer.
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(NSGE-) NS GENE AS.
                                                                                                                                                                                                                                                                                                                                                                                                                          WO200272826-A2.
                                                                                                                                                                                                                                                                                                                                                    Unidentified.
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Babij P, Bex FJ, Yaworsky PJ, Bodine PV;

WPI; 2003-129278/12.

(GENO-) GENOME THERAPEUTICS CORP. (AMHP ) WYETH.

11-MAY-2001; 2001US-0290071P. 17-MAY-2001; 2001US-0291311P. 01-FEB-2002; 2002US-0353058P. 04-MAR-2002; 2002US-0361293P.

13-MAY-2002; 2002WO-US014876

WO200292764-A2

21-NOV-2002

The invention relates to a truncated neublastin polypeptide comprising an amino acid terminus that lacks one or more amino-terminal amino acids of a mature neublastin polypeptide. The polypeptides and nucleic acids are useful for treating neurodegenerative disorders such as ischemic neuronal damage, traumatic brain injury, peripheral neuropathy, neuropathic pain, Alzheimer's disease, Huntington's disease, Parkinson's disease, amyotrophic lateral soletosis, memory impainment, disbetes, real diseases, or glaucoma by moderating metabolism, growth, differentiation or survival of a nerve or neuronal cell. This polynucleotide sequence is a neublastin PCR primer of the invention Sequence 18 BP; 3 A; 7 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ; 3.2%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.3e+02; 2; Indels 0; Mismatches Local Similarity 88.2 ies 15; Conservative Best Loca Matches ð

CCGCGACGACGGCGCCA 396 CIGCGACGACTGCGCCA 18 380

RESULT 180 ACC45873 02-JUN-2003 (first entry)

Human; high bone mass; HBM; LRP5; LRP6; transgenic; bone mass modulation; gene therapy; bone density modulation; bone strangth; trabecular number; bone size; bone tissue connectivity; bone disease; osteoporosis; PCR; osteomalacia; rickets; Paget's disease; neoplasm of the bone; primer; ss.

Homo sapiens

Human HBM STS marker reverse primer #226. ACC45873 standard; DNA; 18 BP ACC45873; g

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The invention relates to novel transgenic animals expressing the high comprising an alteration of the gene encoding LRFS or LRFG, or expressing an alteration of the gene encoding LRFS or LRFG, or expressing an important is modulated by an altered gene control sequence introduced by homologous or non-homologous recombination. The transgenic animals are for the study of bone density modulation or bone mass modulation. The invention has osteopathic and cytostatic activity. The polymucleotides of the invention may have a use in gene therapy. The transgenic animals and condense in more than one parameter selected from bone density, bone cytostatic acids are for the study of bone eissue connectivity. The species in more than one parameter selected from bone density, bone cytostatic acids and methods are useful for identifying molecules involved in bone development, and for developing pharmaceutical compositions, which may be employed for treating or preventing bone connectivity. The compositions, which may be employed for treating or preventing bone connectivity in methods for diagnosing diseases, paget's disease, or neoplasms of the bone. The transgenic animals and nucleic acids are also confositions, when the contained so the parameter sequence is not the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                    New transgenic animals (e.g. mice), useful as models for studying bone density modulation, developing drugs for treating or preventing bone diseases (e.g. osteoporosis), or diagnosing diseases characterized by reduced bone density.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 57; 603pp; English.
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Best Local Similarity 88.2
Matches 15, Conservative
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197 CTGCTCGGTGAAAGCAG 213 crecrassreacascas 17 RESULT 181 ADB98571 ઠે

ADB98571 standard; DNA; 18

ADB98571;

Sequence tagged site #452 used to prepare Zmax1 (LRP5) gene region map. 04-DEC-2003 (first entry) 

Osteopathic, Gene therapy; High Bone Mass; HBM; LRP5; Zmaxl; LRP6; bone mass modulation; osteoporosis; STS; sequence tagged site; ds.

Homo sapiens.

WO200292000-A2

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WPI; 1996-477128/47
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                                                                                                                                                                                                                                                                                             The present invention relates to High Bone Mass (HBM), LRPS (Zmax1) and LRP6 mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bene mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence is a Sequence Tagged Site (STS) marker, which was used to prepare a physical map of the Zmax1 (LRPS) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lymphocyte specific interferon regulatory factor; LSIRF; IRF-3; probe; major histocompatibility complex; MHC; ISRE; interferon-stimulated response element; ds.
                                                                                                                                                                                                            New nucleic acid comprising a mutation in LRP5 or LRP6, useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject
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                                                                                                                                                             Liu W;
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Best Local Similarity 88.2%; Pred. No. 2.38+02;
Matches 15; Conservative 0; Mismatches 2;
                                                                                                                                                            Morales A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Richardson CD;
                                                                                                                                                             Graham JR,
                                                                                                                                                                                                                                                                         Example 2; Page 64; 629pp; English.
                                                                                                                                                                                                                                                   suffering from e.g. osteoporosis.
                                                                                                                      (GENO-) GENOME THERAPEUTICS CORP. (AMHP ) WYETH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      197 CTGCTCGGTGAAAGCAG 213
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                                                         11-MAY-2001; 2001US-0290071F.
17-MAY-2001; 2001US-029131IP.
01-FEB-2002; 2002US-0353058F.
04-MAR-2002; 2002US-0361293P.
                                    13-MAY-2002; 2002WO-US014877
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96US-00611280
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                                                                                                                                                            Allen K, Anisowicz A,
                                                                                                                                                                                     WPI; 2003-129214/12.
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03-APR-1996;
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          21-NOV-2002
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AAT41709
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                                                                                     The murine major histocompability complex interferon-stimulated response element (MHC IRSE) binding sequence (AAT41709) was used as a probe to determine whether novel mouse lymphocyte-specific interferon regulatory factor (LSIRF) (see also AAR99426) is a DNA binding protein. LSIRF polypeptides were incubated with 32P- labelled double-granded probe and, in some cases, with unlabelled competitor DNA fragments (see also AAT41710-16). Gel shift assays showed that the MHC ISRE sequence binds LSIRF protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identification of allele type of a known polymorphic genetic locus - used particularly for human leukocyte antigen allele determination.
genes for murine lymphocyte specific interferon regulatory factor d for modulation of lymphocyte activation and proliferation.
                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphic; Human leukocyte antigen; HLA; DNA sequencing; PCR; polymerase chain reaction; allele; ss.
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                                                                                                                                                                                                                                                                                                                                     3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.68+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                              Sequence 19 BP; 7 A; 2 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 17; 75pp; English.
                                                               Example 4; Page 40; 92pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 CAGAAGTGAAACTGAGG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                               4 CAGGAGTGAAACTGCGG
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Best Local Similarity 88.2
Matches 15; Conservative
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and evaluating four concurrent reactions, the sample is concurrently combined with at most three sequencing reaction mixtures containing different types of chain terminating nucleosides. The method can be used for the evaluation of polymorphic sites, and for determining the allelic type of a polymorphic gene. The methods are particularly useful for determining the HLA allele present in a sample
                                                                                                                                                                                                                                                                                                                                                                            Filamentous flower; FIL protein; agriculture; gardening; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 filamentous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A gene participating in the flower formation of a plant useful in agriculture and gardening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a PCR primer for DNA encoding the filar
flower (FIL) protein of the invention. The protein is useful in
agriculture and gardening
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                                                                                                                    3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; Live 0; Mismatches 2; Indels
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                                                                                            Sequence 19 BP; 2 A; 7 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                    PCR primer for FIL protein coding sequence.
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                                                                                                                                                                                                                                                                      BP
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                                                                                                                                                                           GCCGCGCAGTGGACATC
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                                                                                                                                                                                                                                                                     AAZ49122 standard; DNA; 19
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MITSUI CHEM INC.
DAIICHI ENGEI KK.
TORAY IND INC.
                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                  15; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                       Arabidopsis sp.
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(DAII-) I
(TORA )
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AAC73121/c
ID AAC7311
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AC AAC7311
XX
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Matches
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                                                                                                                                                  Matches
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                                                       Oligonucleotide array; genotyping; single base extension reaction; SBE;
PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pig; muscular steatosis-modulating factor; ss; metabolic; muscular; MS
food supplement; obesity; hyperlipidaemia; atherosclerosis;
wound healing; tumour; amyotrophic lateral sclerosis; ALS; PCR primer.
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                                                                                                                                                                                                                                                                                                                        Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 2 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                          Kaplan P,
                                                                                                                                                                                                                                                                            (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Porcine reverse PCR primer for TGFb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 7; Page 49; 70pp; English.
                                                                                                                                                                                                                                                                                                                          Huang X,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     form the oligonucleotide array
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                                                                                                                                                                                                                               99US-0126473P.
99US-0140359P.
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Ryder T, Sklar P;
                                                                                                                                                                                                                                                                                              (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                      WO200058516-A2.
                                                                                                                                                                                                                               26-MAR-1999;
23-JUN-1999;
                                                                                                         Unidentified.
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Matches
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forward primer #13 used in multiplexing PCR/SBE assay.

(first entry)

02-FEB-2001

(MIAC ) CANADA AGRIC & AGRI-FOOD CANADA. 17-APR-2000; 2000US-0197936P. 12-APR-2001; 2001WO-CA000509 WPI; 2002-017600/02 Homo sapiens Palin M, AAS18013 셤 %#X#X#X#X#X####X#X#X#X#X ઠે

The invention relates to prognosis or diagnosis of muscular steatosis by measuring the level of a muscular steatosis modulating factor (MSMF) in a human or animal and comparing this with the level in a healthy control. Any difference indicates presence of, or predisposition to, muscular steatosis. The method is particularly used for diagnosis or prognosis of muscular steatosis in mammals and birds, e.g. to select individuals as founders in animal breading. Also (ant)agonists of MSMF can be used to treat, or induce (for increasing the fat content of food) muscular steatosis, in humans and animals. The MSMF markers are also useful in the study of diseases and conditions such as obesity, hyperlipidaemia, atherosclerosis, wound healing, tumours and amyotrophic lateral sclerosis invention from its gene ö Prognosis and diagnosis of muscular steatosis, useful e.g. for selecting animals for breeding, by measuring levels of specific markers, also treating or inducing steatosis. Gaps , 0 3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; ative 0; Mismatches 2; Indels Sequence 19 BP; 4 A; 7 C; 6 G; 2 T; 0 U; 0 Other; Example 1; Page 40; 190pp; English. Query Match
Best Local Similarity 88.2 Matches 15; Conservative

Human Neuregulin-2 PCR primer 1531. AAS18013 standard; DNA; 19 BP. (first entry) 12-MAR-2002 AAS18013;

Human; ss; neuregulin-2; NRG-2alpha; NRG-2beta; mitogenesis; cell survival; cell growth; cell differentiation; erbB receptor; cardiomyopathy; ischemic damage; cardiac trama; heart failure; atherosclerosis; vascular lesion; vascular hypertension; 1531; degenerative congenital vascular disease; myasthenia gravis; neurodegenerative disorder; peripheral neuropathy; PCR primer; sensory nerve fiber neuropathy; motor fiber neuropathy; sensory nerve fiber neuropathy; multiple sclerosis; amyotropic lateral sclerosis; spinal mescular atrophy; nerve injury; Alzheimer, s disease; Parkinson's disease; cerebellar ataxia; spinal cord injury; tumour; neurofibromatosis; transgenic animal.

23-MAY-2001; 2001WO-US016896 WO200189568-A1. 29-NOV-2001.

23-MAY-2000; 2000US-0206495P.

(CENE-) CENES PHARM INC.

Marchionni MA;

WPI; 2002-097612/13.

Gariepy C;

Pomar C,

Neuregulin-2 polypeptide and polynucleotide useful for treating multiple sclerosis, spinal muscular atrophy, nerve injury, Alzheimer's disease, by increasing mitogenesis, survival, growth or differentiation of a cell.

Example 1; Page 29; 79pp; English

The invention relates to a substantially pure neuregulin (NRG)-2

COT NRG-2beta (Coince 2b7) and the polymolecideds encoding the Also

COT NRG-2beta (Coince 2b7) and the polymolecideds encoding the Also

included are a vector expressing the protein, a host cell comprising the

corrector, a transgenic non-human animal transformed with the vector or

vector, a walvest of mutation in one or both NRG-2 allelss and an anti-NRG-2

antibody. Analysis of mutations in NRG-2 in an individual is useful for

diagnosing an increased likelihood of developing a NRG-2-related disease

or condition in a test subject. NRG-2 is useful for increasing the

mitogenesis, survival, growth or differentiation of a cell (e.g. a

concomition in a test subject. NRG-2 is useful for increasing the

mitogenesis, survival, growth or differentiation of a cell (e.g. a

concomition in a test subject. NRG-2 is useful for increasing the

concomition in a test subject. NRG-2 is useful for increasing the

concomition in a test subject. NRG-2 is useful for increasing the

concomition in a test subject. NRG-2 is useful for increasing the

concomition in a test subject. NRG-2 is useful for increasing,

and degenerative congenital disease, ischaemic damage, cardiac

conditions and degenerative disease, parkinal muscular arrophy, multiple

conditions and degenerative disease, parkinson's disease, cerebellar ataxia, and

congrishing inhibiting proliferation of a tumour cell, for treating of neurofibromatosis by inhibiting glial cell

congrishing inhibiting of neurofibromatosis by inhibiting glial cell

congrishing inhibiting of neurofibromatosis by inhibiting glial cell

congrishing inhibiting of neurofibromatosis by inhibiting glial cell

congrishing inholessed to serve is a PCR primer used to analyse the

congrishing the present sequence is a PCR primer used to analyse the

Sequence 19 BP; 6 A; 7 C; 5 G; 1 T; 0 U; 0 Other;

Gaps ; 3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; vative 0; Mismatches 2; Indels Query Match Best Local Similarity 88.2 Matches 15; Conservative

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45 GGCCACCACTCAGAGGA

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GGCCACCACACAGACGA 17 RESULT 188

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ABN79916 standard; DNA; 19 BP.

ABN79916,

ABN79916;

Human, single nucleotide polymorphism; nucleic acid typing; primer; tissue typing; PCR; ACE; angiotensin coverting enzyme; ss. Human angiotensin coverting enzyme SNP-fragment Bu6 PCR primer #1. /\*tag= a /note= "Biotinylated" Location/Qualifiers (first entry) Key modified\_base 15-JUL-2002 Homo sapiens CXXXIXEXXXXXXXIIIIXXXX

WO200220837-A2

14-MAR-2002

Pourmand N;

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The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or more nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing three or more variable variable sites are typed, where three or more primer extension reactions are performed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid molecule(s). The methods are particularly suited for identifying microbial species or their subtypes, and in typing procedures e.g. typing of polymorphisms, tissue typing or in clinical applications. The sequence represents a PCR primer used in the invention to amplify a specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adeno-associated virus; AAV; integration locus; CpG island;
SP1-like binding site; CAMP response element; CRE;
upstream binding factor 1; UBF-1; minisatellite; probe; gene therapy;
promoter; amplification; primer; polymerase chain reaction; PCR; ss.
                                                                                                                                                                                                        Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 2 A; 8 C; 4 G; 5 T; 0 U; 0 Other;
                                                                           (PYRO-) PYROSEQUENCING AB.
(STRD ) UNIV LELAND STANFORD JUNIOR.
(GARD/) GARDNER R.
                                                                                                                                                                                                                                                                                            Example 2; Page 47; 86pp; English
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               10-SEP-2001; 2001WO-GB004042.
                                           08-SEP-2000; 2000GB-00022069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ63197 standard; DNA; 20
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nes 15; Conservative
                                                                                                                                          Ronaghi M, Ekstroem B,
                                                                                                                                                                             WPI; 2002-393849/42
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                                                                                                                                                                                                                                                            incorporation.
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18-NOV-1994
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266 GCACCTGGAGCAGGGCG 282

19 GTACCTGGAGCAGAGCG 3

BP.

(revised)
(first entry)

92US-00947127. 93EP-00114941.

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                                                                                                                            In the cloning of AAVS1 from human lung fibroblast DNA, the primers given in AAQ63193-202 were used. A 4kb fragment contg. the AAV integration site was obtained (AAQ63192). (Updated on 25-MAR-2003 to correct PN field.)
                                               New nucleic acid corresponding to human adeno-associated virus integration site - useful e.g., as probe to confirm targetted integration of adeno-associated virus vectors in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New HIV-type immune deficiency virus ECACC V 92092318 - and deriv. cDNA or antigens, useful for diagnosing retroviral infections and vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MVP-5180/91 DNA is obtained by PCR using the primers given in AAQ58925-
958. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-
2003 to correct PI field.)
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retrovirus; vaccine; lymphocyte; reverse transcriptase; amplification;
primer; polymerase chain reaction; PCR; 88.
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                                                                                                                                                                                                             3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; rative 0; Mismatches 2; Indel8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eberle J, Brunn VA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 5; 73pp; German.
Linden RM;
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92DE-04235718.
92DE-04244541.
93DE-04318186.
                                                                                                        Claim 4; Page 4; 20pp; English.
                                                                                                                                                                                                                                                                      81 CGCGCAGTGGACATCAC 97
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(first entry)
                                                                                                                                                                                                                 3.2
Query Match
Best Local Similarity 88.2
Matches 15, Conservative
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 Berns KI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 15; Conserv
                        WPI; 1994-127741/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                tat-1P primer.
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30-DEC-1992;
01-JUN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-OCT-1993;
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04-NOV-1994
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  Kotin RM,
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                                                                                                                                                                                                                                                                                                                                           RESULT 190
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Gaps ;

3.2%; Score 13.8; DB 1; Length 19; 88.2%; Pred. No. 2.6e+02; tive 0; Mismatches 2; Indels

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Synthetic.
Human immunodeficiency virus 1.
                                                                                                  22-OCT-1992;
30-DEC-1992;
01-JUN-1993;
05-OCT-1993;
                                                                                                                                                                   Guertler LG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9963078-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-JUN-1999;
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                                                                        05-OCT-1993;
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                                    EP890642-A2
                                                     13-JAN-1999
                                                                                          16-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Si
Matches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ46578;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ46578
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                            Primers SS01 (given in AAQ76031) and SS02 (AAQ76032) were.used for the PCR amplification of a target region (AAQ76037) in the cytosine-DNA-methyltransferase of N. gonorrhoeae. Probe SS06-T5 (AAQ76033) is specific for a region in the amplified sequence, and is used to identify N. gonorrhoeae. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-type retrovirus, MVP-5180/91; BCACC V 92092318; antigen, assay kit; detection; antibody; immune deficiency; vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                Detection of Neisseria gonorrhoeae and/or Chlamydia trachomatis simultaneously by a simple, rapid and sensitive technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 3.2%; Score 13.8; DB 1; Length 20; Best Local Similarity 88.2%; Pred. No. 2.9e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 20 BP; 5 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                         Neisseria gonorrhoeae, probe, hybridization, cytosine-DNA-methyltransferase, CMT, ss.
                                                                                                                                                                                                                                                                                          (HOFF ) HOFFMANN LA ROCHE & CO AG F.
                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Fig 1; 29pp; English.
         240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      367 TCACTTTCCTGGACCGC 383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX22342 standard; DNA; 20 BP
                                                                       AAQ76033 standard; DNA; 20 BP.
                   17 GGATGCTTCCAGGGCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      н
                                                                                                                                        N. gonorrhoeae probe SS06-T5.
                                                                                                                                                                                                                                             94EP-00108997
                                                                                                                                                                                                                                                              93US-00082851
94US-00214861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 TCACTTCCCTGAACCGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1 PCR primer tat 1P.
                                                                                                                                                                                                                                                                                                            Purohit AP, Silver SB;
                                                                                                             (revised)
                                                                                                                                                                                                                                                                                                                               WPI; 1995-031607/05.
                                                                                                                                                                                                                                             13-JUN-1994;
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19-MAY-1999
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                                                                                                             25-MAR-2003
16-JUL-1995
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AAX22342/
                                                                AAQ76033
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This invention describes the isolation of a novel HIV-type retrovirus called MVP-5180/91 (ECACC V 92092318). Antigens produced from this product can be used in an assay kit for detecting antibodies against viruses that cause immune deficiency, preferably where the assay is a Western blot, ELISA or fluorescence immunoassay. MVP-5180/91, cDNA and/or antigen can be used for detecting retroviruses that cause immune deficiency and to prepare vaccines. This sequence represents a PCR primer used in the method of the invantion. (Updated on 20-MAR-2003 to correct PF field.) (Updated on 20-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Retinal calcium channel; RCC gene; alphalF-subunit; retinal disorder; myopia; nystagmus; strabismus; calcium-regulated development pathway; eye disorder; human; CACNAIF; CSNB; mutational analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                       New HIV-type retrovirus and corresponding cDNA, recombinant DNA and antigen - used for detecting retro-viruses that cause immune deficiency and to prepare vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                     Hauser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Forward primer specific for human CACNALF exon 16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.2%; Score 13.8; DB 1;
88.2%; Pred. No. 2.9e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                              Eberle J, Brunn AV, Knapp S,
                                                                                                                                                                                                                                                     (DADE-) DADE BEHRING MARBURG GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 4; 39pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              256
                                                                                    92DE-04235718.
92DE-04244541.
93DE-04318186.
93EP-00116058.
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98EP-00114623
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ46578 standard; DNA; 20
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hes 15; Conservative
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rng.res

Disclosure, Page 78; 115pp; English

(UYTE-) UNIV TECHNOLOGIES INT INC. Bech-Hansen T, Naylor MJ;

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New isolated mammalian retinal calcium channel gene, used to develop products for the diagnosis and treatment of incomplete congenital stationary night blindness and related disorders.
                                                                                                                                                                                                                                                             Disclosure, Fig 6; 55pp; English
WPI; 2000-097327/08.
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The invention provides a DNA molecule comprising a sequence of nucleotides encoding an alphalF-subunit of a mammalian retinal calcium channel (RCC), including a human alphalF-subunit, a murine alphalF-subunit and orthologs of the human and murine alphalF-subunits. The RCC gene may be used to develop products for diagnostic tests, for incomplete CSMB and risk assessment in affected families. The RCC gene can provide information as to the basic defect in this retinal conditions, which could lead to effective methods for treatment or cure of the disorder. As the associated features of myopia, nystagmus and strabismus frequently observed in patients with incomplete CSMB may be caused by calcium-requiated development pathways, identification of the RCC gene may help to elucidate the molecular details of eye development and which may lead to reatment for related eye disorders or diseases. Sequences AAZ46563-650 represent human CACNALF (alphalF-subunit of RCC gene) exon-specific PCR primers, used for mutational analysis in humans

Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

Gaps .; 0 Match 3.2%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 2.9e+02; es 15; Conservative 0; Mismatches 2; Indels Query Match Matches

ઠે 9 AAZ44577 standard; DNA; 20 BP

AAZ44577;

Newcastle disease virus LaSota primer p1898-.

07-APR-2000 (first entry)

Avian-paramyxovirus; infection; lentogenic; F protein; vaccine; respiratory disease; gastrointestinal disease; poultry pathogen; local immunity; primer; ss. 

Newcastle disease virus

WO9966045-A1.

23-DEC-1999,

99WO-NL000377 .7-JUN-1999; DIEN-) STICHTING DIENST LANDBOUWKUNDIG ONDERZOE.

98EP-00202054

19-JUN-1998;

Gielkens ALJ; De Leeuw OS, Koch G, Peeters BPH,

WPI; 2000-106102/09.

New avian paramyxovirus cDNA, useful for production of vaccine against Newcastle disease virus.

This invention describes a novel avian-paramyxovirus cDNA (I) which comprises a nucleic acid sequence corresponding to the 5' terminal end of the genome of avian-paramyxovirus allowing the generation of an infectious copy of avian-paramyxovirus. The cell line is useful for the production of infectious lentogenic NDV (Newcastle Disease virus) without the addition of exagences proteolytic activity. Also it is possible to generate a stable transfected cell line that expresses the wild-type F protein in the virus envelope therefore providing infectious particles, useful in the form of a vaccine, especially adainst respiratory and/or gastrointestinal diseases. NDV can be easily cultured to very high titers in embryonated eggs. Mass culture of embryonated eggs is relatively chapte as application methods e.g. drinking water or by spraying or by across are relatively stable and can be simply administered chap. Mass application methods e.g. drinking water or by spraying or by across prointestinal tract which are also the major routes of infection of many other poultry pathogens. NDV can induce local immunity despite the presence of circulating maternal antibody. AAZ44618-244650 represent primers used in the isolation of the NDV strain LaSota genome 

Sequence 20 BP; 4 A; 4 C; 6 G; 6 T; 0 U; 0 Other;

Gaps ö / Match 13.8; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 2.9e+02; Nes 15; Conservative 0; Mismatches 2; Indels Query Match Best Local S: Matches 15

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C-terminal phenylalanyl-tRNA synthetase DNA amplifying primer, BfP-5.

Phenylalanyl-tRNA synthetase; PheRS; amino acid separation; ATP quantitation; protein inhibitor; antimicrobial; antibiotic effect; PCR primer; ss.

97US-00855910.

14-MAY-1997; 97US-00855910.

Avruch AS, Shen X, Sassanfar M, Gallant PL,

Yu RV;

New Enterococcus faecalis aminoacyl-tRNA synthetase proteins and nucleic acids useful for separating amino acids which they specifically recognize, in quantifying amino acids and ATP, or for detecting protein inhibitors. 

Example 3; Col 41; 88pp; English.

The present invention relates to Enterococcus faecalis aminoacyl-tRNA synthetases are useful in the biochemical

Page

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geparation of the amino acid which they specifically recognise and in quantitations of the amino acid and ATP, and for detecting and indentifying inhibitors of their activities. The potential inhibitors of these enzymes can be screened for antimicrobial or antibiotic effects, without requiring the culture of pathogenic strains of Enterococcus. The antibodies which bind to these enzymes can be made and used in the purification and study of the enzymes. The aminoacyl-tRNA synthetase genes may be used as probes to identify DNA fragments are used in the production of proteins or polypeptides, and the aminoacyl-tRNA synthetase genes may be used as probes to identify DNA fragments encoding the corresponding aminoacyl-tRNA synthetase gene from other species of enterococci by specific hybridisation. The present sequence is a PCR primer which is used for amplifying the C-terminal Enterococcus faecalis phenylalanyl-tRNA synthetase (PheRS) DNA
              88888888888888888
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Sequence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;

. 0 3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; ative 0; Mismatches 2; Indels Query Match
Best Local Similarity 88.2
Matches 15; Conservative

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RESULT 196 AAS09653/C 

AAS09653 standard; DNA; 20 BP. AAS09653; (first entry) 26-SEP-2001

Immunoreactive CpG sequence-containing oligonucleotide #103.

CpG sequence; immune response; non-B cell activation; interferon gamma; IFN-gamma; humoral; antibody production; interleukin-6 production; therapeutic; allergy; asthma; cancer; autoimmune disorder; infection; bio-warfare; vaccine; antisense therapy; eczema; allergic rhinitis; coryza; hay fever; urticaria, hives; food allergy; atopic condition; hepatitis; human immunodeficiency virus; HTV, malaria; Francisella; lupus erythematosus; rheumatoid arthritis; multiple sclerosis; schistosomiasis; tuberculosis; acquired immunodeficiency syndrome; AIDS; leishmania; Ebola; Anthrax; Listeria; ss.

Synthetic

WO200151500-A1.

19-JUL-2001

12-JAN-2001; 2001WO-US001122

14-JAN-2000; 2000US-0176115P.

(USSH ) US DEPT HEALTH & HUMAN SERVICES.

Verthelyi D; Ishii K, Klinman D,

WPI; 2001-442129/47.

Oligodeoxynucleotides for inducing an immune response to treat and prevent an allergic reaction, cancer, an autoimmune disorder and symptoms resulting from exposure to bio-warfare agents, comprise multiple CpG sednences

Claim 5; Page 44; 48pp; English.

AASO9551-AASO9662 represent oligodeoxynuclectides (ODN) of at least 10 nuclectides comprising multiple CpG sequences, where one of the CpG sequences is different from another of the multiple CpG sequences. The ODN are useful for inducing an immune response, preferably a cell-mediated immune response, preferably a cell-mediated immune response, involving non-B cell activation, interferon

cell activation, antibody and interleukin-6 production in a host, for treating, prevention, antibody and interleukin-6 production in a host, for treating, preventing or ameliocating an allergic reaction, e.g. asthma, cancer, e.g. solid tumour cancer, a disease associated with the immune cystem e.g. autoimmune disorder or an immune system deficiency, infection of a symptom resulting from exposure to bio-warfare agent in a human. The corduction of immune response improves the efficacy of a vaccine and isc used in antisense therapy. The ODN are useful for treating, preventing or used in antisense therapy. The ODN are useful for treating, preventing or coryza, hay fever, bronchial asthma, urticaria (hives), food allergic radictions, including eczema, allergic radictions, for improving the efficacy of vaccines or adainst hepatitis A, B and C, human immunodeficiency virus (HIV) and malaria, for treating immune system deficiencies, e.g. lupus corythematosus and autoimmune diseases such as rheumatoid arthritis and comultiple sclerosis, infections including Francisealla, eschistosomiasis, cuberculosis, acquired immunodeficiency syndrome (AIDS), Leishmania and symptoms resulting from exposure of bio-warfare agent, including Ebola, xxx

Sequence 20 BP; 2 A; 4 C; 12 G; 2 T; 0 U; 0 Other;

Gaps ö 3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; ive 0; Mismatches 2; Indels Query Match
Best Local Similarity 88.4.
Local 15; Conservative

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RESULT 197 AAS21720/c

B AAS21720 standard; DNA; 20

(first entry) 21-NOV-2001 AAS21720; 

Mouse Survivin antisense oligonucleotide #23.

Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis;

Mus musculus. Synthetic WO200157059-A1.

09-AUG-2001.

30-JAN-2001; 2001WO-US002939.

02-FEB-2000; 2000US-00496694.

(ISIS-) ISIS PHARM INC

Cowsert LM; Swayze EE, CF, Ackermann EJ, Bennett

WPI; 2001-488863/53.

Novel antisense compounds for modulating the expression of Survivin and treatment of cancer.

Example 18; Page 60; 120pp; English.

The invention relates to antisense oligonucleotides targeted to a nucleic acid molecule encoding human Survivin, where the antisense oligonucleotide inhibits the expression of human Survivin. These antisense oligonucleotides are used in the treatment of an animal suffering from a disease or condition associated with Survivin, e.g. a hyperproliferative condition such as cancer, and comprises administering a therapeutically or prophylactically effective amount of the antisense oligonucleotide so that expression of Survivin is inhibited. The

Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis; signal transduction; DNA replication; cell division; growth; proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.

Candida albicans. WO200253728-A2.

Candida albicans GRACE strain PCR primer SEQ ID NO 4516.

(first entry)

30-JAN-2003

ABZ30365;

ABZ30365 standard; DNA; 20 BP.

RESULT 199

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oligonucleotides can also be used to treat a human suffering from a disease or condition characterised by a reduction in apoptosis comprising administering the antisense oligonucleotide to a human. In addition, the antisense oligonucleotide and a cytotoxic chemcherapeutic agent e.g. taxol or cisplatin, can be used to modulate apoptosis, cytokinesis or the cell cycle, or inhibit the proliferation in a cancer cell by contacting the cell with the antisense oligonucleotide. AAS1521-AAS21768 represent Survivin nucleic acids, and antisense oligonucleotides targeted to Survivin, used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides a microarray of oligonuclectides comprising probes for the human HLA Class I genes attached to a solid support. These can be used in HLA typing. Oligonuclectide arrays are also useful in large scale gene discovery, monitoring gene expression, polymorphism detection and gene mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide arrays useful for human leukocyte antigen (HLA) tissue
typing, comprises HLA class I oligonucleotide probes representing all
known polymorphisms in HLA class I locus, on a solid support.
                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, HLA typing, oligonucleotide array; Class I; gene discovery; expression; polymorphism detection; mapping; probe; PCR primer; ss.
                                                                                                                                                                                                                             3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human HLA Class I oligonuclectide probe SEQ ID NO: 38.
                                                                                                                                                                                      Sequence 20 BP; 6 A; 3 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUTC-) HUTCHINSON CANCER RES CENT FRED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 54; 83pp; English.
                                                                                                                                                                                                                                                                                                     358 GCGACTICCTCACTITC 374
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF54593 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guo Z,
                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 15, Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-102734/11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF54593;
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The invention relates to constructing (M1) a strain of diploid fungal calls in which both alleles of a gene are modified, comprising modifying calls in which both alleles of a gene are modified, comprising modifying cone allele by insertion or replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells in which both alleles of a gene are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a fungus; a gene chart, an antifungal agent that inhibits the growth of a mammalian disease. (M1) is useful for identifying a compound which modulates the compound catabolism, biosynthetic, transporter, transcriptional, transduction, part is compound having the ability to inhibit growth or proliferation of C. albicans cells and for transtraint infection by C. albicans. The present sequence is that of a por primer used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ohlsen KL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bussey H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.2%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roemer T, Jiang B, Boone C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           228 GCCAAATCGGGAGGCTG 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GCCAAATCGGAAGACTG 17
                                                                                                                                                                                                                                                                                                                                      26-DEC-2001; 2001WO-US049486.
                                                                                                                                                                                                                                                                                                                                                                                                 20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
                                                                                                                                                                                                                                                                                                                                                                              29-DEC-2000; 2000US-0259128P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 88.2
les 15; Conservative
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Gaps

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2; Indels

0; Mismatches

298 AGGACCTGAGCCCCGGG 314

15; Conservative

Matches

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Similarity

Query Match Local AGGACCTGAGCTCCTGG 18

3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02;

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The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying cells in which both alleles of a gene are modified, comprising modifying one allale by insertion or replacement by asserte having an expressible selectable marker and modifying other allele by recombination, of a promoter replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that is essential to the survival or growth of a fungus, a gene that contributes to the relatione and/or pathogenicity of a fungus, a gene that contributes to the relational of a diploid fungus to an antifungal agent that inhibits the growth of a mammalian disease. (M1) is useful for identifying a compound extabolism, biosynthetic agent for treatment of a mammalian disease. (M1) is useful for identifying a compound having the compound actabolism, biosynthetic, transporter, transcriptional, transcribly ensymatic activity, carbon compound actabolism, biosynthetic, transporter, transcriptional, transcrible growth or proliferation of C. albicans cells and for treatment is not represented in the princed specification but is based on sequence information supplied to Derwent by the Buropean Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                          Fungus; yeast; tetracyclin; promoter; GRACE strain; biosynthesis;
signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 36; SEQ ID NO 5310; 167pp + Sequence Listing; English.
                                                                                                                                                                    Candida albicans GRACE strain PCR primer SEQ ID NO 5310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bussey H, Ohlsen KL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 7 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
                                     ABZ31091 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boone C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
                                                                                                                                                                                                                                                                                                                                                                                                                                     26-DEC-2001; 2001WO-US049486.
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                                                                                                                          30-JAN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roemer T, Jiang B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-566694/60.
                                                                                                                                                                                                                                                                                                   Candida albicans.
                                                                                                                                                                                                                                                                                                                                              WO200253728-A2.
                                                                                                                                                                                                                                                                                                                                                                                          11-JUL-2002.
                                                                                  ABZ31091;
RESULT 200
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3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; ative 0; Mismatches 2; Indele Query Match Best Local Similarity 88.2 Matches 15; Conservative

1 GCCAAATCGGAAGACTG 17 임

AAD45182,

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Human, receptor interacting protein, RIP2; antisense; gene therapy; phosphorothicate; ss.
                                                                                                                                        note= "2-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                         note = "2-methoxyethyl (2'-MOE) nucleotides"
                                                                                                              note= "Phosphorothioate backbone"
                                     Human RIP2 antisense oligonucleotide ISIS #104252.
                                                                                       Location/Qualifiers
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/*tag= a
/mod_base= OTHER
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/mod_base= OTHER
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AAD45182 standard; DNA; 20 BP
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                                                                   Homo sapiens.
Synthetic.
            AAD45182;
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US6426221-B1

30-JUL-2002.

01-AUG-2001; 2001US-00920663.

01-AUG-2001; 2001US-00920663.

(ISIS-) ISIS PHARM INC.

Ward DT, Cowsert LM;

WPI; 2002-673017/72.

New antisense oligonucleotide that targets regions of a nucleic acid encoding human receptor interacting protein (RIP)2, for treating diseases associated with RIP2 expression.

Claim 3; Col 46; 35pp; English.

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0; Gabs

The invention relates to antisense compounds targetted to a nucleic acid encoding human receptor interacting protein (RIP)2 to inhibit its expression. Antisense compounds are used for treating diseases associated with RIP2 expression. They are also useful in antisense gene therapy. The present sequence is an oligonucleotide targetted to human RIP2 DNA

228 GCCAAATCGGGAGGCTG 244

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Best Local Similarity
Matches 15; Conserv
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                                                                                                    RESULT 203
ABI94844/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel isolated nucleic acid molecule comprising a sequence that encodes a thioesterase or thioesterase domain, derived from a bacterial daptomychin biosynthetic gene cluster. The proteins of the invention have antibacterial, fungicide, virucide, antiparasitic, immunomodulator, antilipemic, and cytostatic activity. The polynucleotides may have a use in gene therapy. The compositions and methods of the present invention are useful for generating novel linear and cyclic peptides and improving yield of a product in a cell expressing and appromych non-ribosomal peptide synthetase (NRES) to be used as new compounds or in producing new compounds, such as antibiotics, antitumour agents, antifungals, anti-cholesterolemic agents, anti-cholesterolemic agents, affectophores, approximately agents, anti-cholesterolemic agents, incompounds and cycostatics. The sequence represents a PCR primer used in the invention to amplify the S. roseosporus daptomycin biosynthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated nucleic acid molecule from a bacterial daptomycin biosynthetic gene cluster encoding a thioesterase or thioesterase domain, useful for generating novel linear and cyclic peptides, and products in a cell.
                                                                                                                                                                                                                                                                                 Daptomycin biosynthetic gene cluster; thioesterase; antibacterial; fungicide; virucide; antiparasitic; immunomodulator; antilipemic; cytostatic; gene therapy; antimitotic; immunomodulatory; siderophore; anti-cholesterolemic; agrochemical; linker; PCR; primer; ss.
                                                                   Gaps
                                                                                                                                                                                                                                                          S. roseosporus daptomycin biosynthetic gene cluster PCR primer P76
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                                     / Match 3.2%; Score 13.8; DB 1; Length 20; Local Similarity 88.2%; Pred. No. 2.9e+02; les 15; Conservative 0; Mismatches 2; Indels
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              Sequence 20 BP; 2 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Silva CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 91; 227pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene cluster from a BAČ library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baltz RH,
                                                                                             318
                                                                                                                                                                                 ABQ78909 standard; DNA; 20 BP
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28-FEB-2001; 2001US-0272207P.
06-AUG-2001; 2001US-0310385P.
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                                                                                                                    ccreaccccacacc 4
                                                                                             CCTGAGCCCCGGGGACC
                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                   Streptomyces roseosporus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Miao VPW, Brian P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-599794/64
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BALTZ R H.
SILVA C J.
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                                                                                                                                                                                                                                   23-OCT-2002
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                                                                                                                                                                                                            ABQ78909;
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(BALT/)
(SILV/)
                                           Query Match
                                                                                                                                                           RESULT 202
                                                                   Matches
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Length 20;

DB 1;

3.2%; Score 13.8;

Query Match

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Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with agene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forenaics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and infrared microscope) the support at the set occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABIS2074 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method (M1) for designing capture oligonuclectide probes (I) for use on a support to which complementary oligonuclectide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus funnigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and pollo virus, and parasitic infectious agents medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human, K-ras, PCR primer, probe, capture probe, mutation detection, ligase detection reaction, LDR, p53, BRCA1, BRCA2, infectious disease, infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer, oncogene, tumour suppressor, human papillomavirus, forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                  Gaps
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                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
88.2%; Pred. No. 2.9e+02;
ive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Capture oligonucleptide Zip ID#1931 oligo #9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Fig 29; 300pp; English
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                                                                                                          373 TCCTGGACCGCGACGAC 389
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                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                  TACTGGACCTCGACGAC
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                                      15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-034366/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABI94844;
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dystunction and a second active agent comprising an entilnflammatory steroid and ubjquinone. A composition may have a mutinflammatory steroid and ubjquinone, A composition may have a immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an preventing sensitivity to adenosine, reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubjquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed of specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine seceptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                         Gaps
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                         Indels
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  Best Local Similarity 88.2%; Fred. No. 2.9e+02;
Matches 15; Conservative 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ ID NO 447; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                  361
                                                                                                                                                                                                                                                                                                                                                   Human oligonucleotide sequence.
                                                                                                                                                                                                                   BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                ABZ85205/c
ID ABZ85205 standard; DNA; 20
                                                                  345 CGGCTGCTCTACAGCGA
                                                                                                         18 CGGCTGCGATACAGCGA
                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-OCT-2002
                                                                                                                                                                                                                                                                                                        17-0CT-2003
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Miller S,
                                                                                                                                                                                                                                                                 ABZ85205;
                                                                                                                                                                            RESULT 204
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The present invention describes a method and a kit for determining the expression of mRNA or cDNA of a protein participating in the maintenance of skin structure. The method is quantitative, simple and accurate in the determination of extracellular matrix components of laminin 5 chain genes LAMA3, LAMB3 and LAMC2, matrix metalloproteinases MMP-1, MMP-2, MMP-3 and MMP-9, VII collagen, type I collagen alpha 1 chain, type I collagen alpha 1 chain, type IV collagen alpha 1 chain, type IV collagen alpha 2 chain, TIMP-1, TMMP-3 and TIMP-3. ACF57290 represent PCR primers and probes used in the method of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; mouse; skin structure; skin; laminin 5 chain gene; LAWA3; LAMB3; LAMC2; extracellular matrix component; matrix metalloproteinase; MMP-1; MMP-2; MMP-3; MMP-3; MMP-9; TIMP-1; TIMP-2; TIMP-3; collagen; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A method and a kit for determination of expression of mRNA or cDNA of protein participating in the maintenance of skin structure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                       Gaps
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                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 2 A; 4 C; 8 G; 6 T; 0 U; 0 Other;
88.2%; Pred. No. 2.9e+02;
ive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                            Human TIMP-2 reverse PCR primer SEQ ID NO:83.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 4; 34pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    144 GCGGTGGAGGCCGGCTT 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
                                                                      245 CTTCCCGGGCTCGGCCA 261
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD56986 standard; DNA; 20
                                                                                                                           CTACCAGGCTCGGCCA
                                                                                                                                                                                                                                                 ACF57283 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15; Conservative
                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SHIS ) SHISEIDO CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-407328/39.
                                                                                                                                                                                                                                                                                                                                                  (first
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JP2002330792-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                                                                                                                                                                                ACF57283;
                                                                                                                           38
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                                                                                                                                                                                                    RESULT 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD56986
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Length 20;

DB 1;

Score 13.8;

Query Match

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New compound, having a sequence targeted to a nucleic acid encoding mucin 1, transmembrane, useful for preparing a composition for treating hyperproliferative or inflammatory disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to antisense oligonuclectides targetted to a nucleic acid encoding mucin 1 transmembrane (also known as MUC1, epistalin, epiterctin, polymorphic epithalial mucin; PBM, peanut-reactive urinary mucin; PDM, epithalial membrane antigen; BMA, PBS-0, NCRC11, H23 antigen, H23-ETA transmembrane antigen, DF3 antigen and CD227) to inhibit/modulate the expression of mucin 1 transmembrane. Antisense compounds of the invention are useful for preparing compositions for treating hyperproliferative or inflammatory disorders. The invention is also used in gene therapy. The present sequence is human mucin 1 transmembrane antisense oligonuclectide
          Human, mucin 1 transmembrane, hyperproliferative disorder; cytostatic, inflammatory disorder; gene therapy, H23-ETA transmembrane antigen; antisense; episialin; epitectin; polymorphic epithelial mucin; CD227; peanut-reactive urinary mucin; PUM; epithelial membrane antigen; BEA; PEM; NCRC11; H23 antigen; DF3 antigen; phosphorothicate backbone; MUC1;
                                                                                                                                                                                                       /mod_base= OTHER
/note= "Phosphorothioate backbone; All cytidines are 5-
methyl cytidines"
                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "2'-methoxyethoxy (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                  mod_base= OTHER
'note= "2'-methoxyethoxy (2'-MOE) nucleotides"
16. .20
                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 82; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-DEC-2001; 2001US-00029517.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-DEC-2002; 2002WO-US039873
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                                                                                                                                                                                                                                                                     ...5
*tag=
                                                                                                                                                                                                                                                                                                                                                   tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Myers SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-559135/52.
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                                                                                                                                                                        Key
modified_base
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                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                       sapiens.
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                                                                                         PAS-0, ss
                                                                                                                                       Synthetic
                                                                                                                         Ношо
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Gaps . 3.2%; Score 13.8; DB 1; Length 20; 88.2%; Pred. No. 2.9e+02; ive 0; Mismatches 2; Indels Sequence 20 BP; 4 A; 3 C; 6 G; 7 T; 0 U; 0 Other; Query Match Best Local Similarity 88.2 Matches 15; Conservative

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247
                   20
231 AAATCGGGAGGCTGCTT
                    ATATCGAGAGGCTGCTT
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B5. ADB89961 standard; DNA; 20 ADB89961 SX S

RESULT 207

ADB89961;

04-DEC-2003 (first entry)

Antinsense oligonucleotide targeting mouse C3 component, ISIS140049

Mouse; ss; antisense; complement component C3; inflammation; septic shock; multiple organ failure; hyperacute organ failure; autoimmune disorder. CNS inflammation; multiple sclerosis; tumour.

Mus musculus

Location/Qualifiers Д Key modified\_base

 $\text{note}_{=}$  "Phosphorothioate backbone and all cytosines are methyl cytosines" mod\_base= OTHER modified\_base

\*tag≕ a

/mod\_base= OTHER /note= "2'-methoxyethyl nucleotides" 16. .20 /mod\_base= OTHER /note= "2'-methoxyethyl nucleotides" υ \*tag= modified base

US2003096775-A1

22-MAY-2003.

23-OCT-2001; 2001US-00001076.

23-OCT-2001; 2001US-00001076.

(ISIS-) ISIS PHARM INC

Watt AT; Graham MJ,

WPI; 2003-606441/57.

New antisense oligonucleotides targeted to a nucleic acid molecule encoding complement component C3, useful for treating a disease or condition associated with complement component C3, e.g. autoimmune disorder or infection.

Claim 3; Page 27; 72pp; English.

The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding complement component C3. The compound complement component C3 and inhibite the expression of complement component C3 and inhibite the expression of complement component C3. Or specifically hybridises with at least an 8-nucleobase component C3. Also included are a composition of complement component C3. Also included are a composition comprising complement component C3. Also included are a composition comprising complement component C3 in calls or tissues (comprising the expression of complement component C3 in calls or tissues (comprising contacting the cclis or tissues with the compound cited above) and treating an animal captures and administering to the animal the compound cited above so that expression of complement component C3 comprising administering to the animal the compound cited above so that compounds are useful for inhibiting the expression of complement c3 in cells or tissues, or for treating an animal having a disease or condition associated with complement component C3 such as an caucimmume disorder (e.g. multiple sclerosis), an infection, or atherosclerosis, inflammation, septic shock, multiple organ failure, computed the capture sclerosis, inflammation, septic shock, multiple organ failure and CNS inflammation. The compounding are also custful as research reagents and diagnostics, in distinguishing functions of various members of a biological pathway, or for preventing or delaying inflammation, inflammation. The present sequence is an animal parties and parties and path an antisense oligonucleotide targeting mouse C3. 

Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;

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Gaps

; 0

Length 20;

Score 13.8; DB 1; Length 2 Pred. No. 2.9e+02; ); Mismatches 2; Indels

; 0

3.2%;

3.2 Query Match Best Local Similarity 88.2 Matches 15; Conservative

254 CTCGGCCACGGTGCACC 270

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0

cccreccaceerecace 1

17

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The present invention describes a method for inducing the production of vascular endothelial growth factor (VEGF) by a cell comprising contacting the cell with a CpG oligonucleotide and therefore inducing the production of VEGF by the cell. Also described: (1) inducing neovascularisation in a tissue, comprising introducing a CpG oligonucleotide into an area of the tissue, where the formation of new blood vessels is desired, and so inducing neovascularisation in the area of the tissue; (2) promoting angiogenesis in an area of the subject where angiogenesis is desired, comprising introducing a CpG oligonucleotide to the area, and so promoting angiogenesis in the subject; and (3) screening for an agent that inhibits neovascularisation, comprising administering a CpG oligonucleotide to a non-human mammal and administering a CpG oligonucleotide to a non-human mammal and administering the agent to the agent is effective in inhibiting neovascularisation. The CpG oligonucleotides have vulnerary, vasotropic and antiarteriosclerotic capett is effective in inhibiting neovascularisation. The method and the CpG oligonucleotides can be used in gene therapy. The method and the CpG oligonucleotides can be used in new thord and such the capette in each as in subjects with a skin graft, subjects who exhibit male pattern baldness, or subjects with a skin graft, subjects who catherents at inhibit neovascularisation. The present sequence the present invention of the present invention of the present sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inducing the production of vascular endothelial growth factor by a cell, useful for inducing angiogenesis, comprises contacting the cell with a CpG oligodeoxynucleotide.
                                                                                                                                                                                                                                                                                                                                                                           vascular endothelial growth factor; VEGF; CpG oligonucleotide;
neovascularisation; angiogenesis; vulnerary; vasotropic;
antiarteriosclerotic; gene therapy; skin graft; male pattern baldness;
atherosclerosis; ischaemia; ss.
                                         Gaps
                                         ;
0
Score 13.8; DB 1; Length 20;
Pred. No. 2.9e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYTE-) UNIV TENNESSEE RES CORP.
(USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 7; SEQ ID NO 45; 37pp; English.
                                                                                                                                                                                                                                                                                                                                          CpG D oligonucleotide SEQ ID NO:45.
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                                                                                                                       20
                                                                                 54
                                                                                                                                                                                                                       ADD01081 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-DEC-2001; 2001US-0343457P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-DEC-2002; 2002WO-US040955.
    3.2%;
                                                                                 38 CGAAGATGGCCACCT
                                                                                                                       CGAAGTTTGCCACCACT
                                                                                                                                                                                                                                                                                                  (first entry)
                         Similarity 88.2
5, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zheng M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-559138/52.
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                                                                                                                                                                                                                                                                                                  01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Klinman DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JUL-2003
                                           15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                               ADD01081;
      Query Match
Best Local
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The sense primer B3', tther with antisense primer B4' (see AAT16479), can be used in the polymerase chain reaction for amplification of the primate alpha-herpes virus gB glycoprotein gene in clinical or laboratory specimens. Following digestion of the amplified product with a restriction endonuclease (e.g. HaeIII), which is not capable of digesting herpes simplex virus (HSV)-1 and HSV-2, the digested fragments may be separated by size or may be hybridized with end-labelled oligonucleotide probe [B8].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 2 A; 5 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.2%; Score 13.8; DB 1;
88.2%; Pred. No. 3.2e+02;
iive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22 TGACCGAGGGCTGGGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 88.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-SEP-1996
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Best Local S
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Sequence 20 BP; 2 A; 4 C; 12 G; 2 T; 0 U; 0 Other;

Detection of herpes B virus by PCR amplification of sample DNA - to detect a specific herpes simian monkey B virus DNA segment.

Claim 4; Col 35; 22pp; English.

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Black

Eberle R,

Scinicariello F,

Hilliard J,

WPI; 1996-105220/11.

(SWBI-) SOUTHWEST FOUND BIOMEDICAL RES.

93US-00042747.

93US-00042747.

01-APR-1993; 01-APR-1993;

30-JAN-1996.

US5487969-A Synthetic.

Primer; polymerase chain reaction; PCR; diagnosis; herpes B virus; primate alpha-herpes virus gB glycoprotein; ss.

B3' for primate alpha-herpes gB glycoprotein.

Sense primer

(first entry)

11-MAY-1996

AAT16477;

BP.

AAT16477 standard; DNA; 21

RESULT 209

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Wed Apr 21 12:58:21 2004
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The present sequence is an antisense oligomucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The oligomucleotide has an average EC(90) (nM) of 150o, which refers to the comc. of oligomucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothioate linkages only)
                                                                                                                                                                                                                                                                                       Oligo:nucleotide(s) corresponding to HIV sequences - used for the detection of HIV or for inhibiting HIV propagation, partic. in infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus; HIV; antisense oligonucleotide; tat;
detection; treatment; infection; inhibition; p24; core antigen;
production; ss.
                       Human immunodeficiency virus; HIV; antisense oligonucleotide; tat; detection; treatment; infection; inhibition; p24; core antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     / Match 3.2%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 3.2e+02; nes 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 2 A; 7 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV tat targetting antisense oligonucleotide.
tat targetting antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                              Example 3; Page 50; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT32083 standard; RNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94US-00277857
                                                                                                                                                           95WO-US009080.
                                                                                                                                                                                   94US-00277857
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                                                                                                                                                                                                              (GENP-) GEN-PROBE INC
                                                                                                                                                                                                                                      Ryder TB, Kwoh TJ;
                                                                                                                                                                                                                                                                WPI; 1996-105849/11
                                                     production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9602557-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUL-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L6-SEP-1996
                                                                                                                                                           14-JUL-1995;
                                                                                                                                                                                     19-JUL-1994;
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                                                                                                       WO9602557-A1
                                                                                                                                  01-FEB-1996
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                                                                              Synthetic.
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                                                                                                                                                                                                                                                                                                                      subjects.
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  HIV
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Kwoh TJ;

Ryder TB,

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The present sequence is an oligomucleotide complementary to an antisense oligomucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The DNA equivalent of the antisense oligomucleotide has an average EC(90) (mM) of 1500, which refers to the conc. of oligomucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothicate
                                                                                                                                      The present sequence is an antisense oligonucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The DNA equivalent of the oligonucleotide has an average EC(90) (LMV of 1500, which refers to the conc. of oligonucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothicate linkages only)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligo:nucleotide(s) corresponding to HIV sequences - used for the detection of HIV or for inhibiting HIV propagation, partic. in infected
                                         Oligo:nucleotide(s) corresponding to HIV sequences - used for the detection of HIV or for inhibiting HIV propagation, partic. in infected
                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide complementary to HIV tat targetting antisense oligo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus; HIV; antisense oligonucleotide; tedetection; treatment; infection; inhibition; p24; core antigen; production; complementary; ss.
                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 64.7%; Pred. No. 3.2e+02;
Matches 11; Conservative 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 6 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                       Sequence 21 BP; 2 A; 7 C; 6 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Page 69; 90pp; English.
                                                                                                            Example 3; Page 56; 90pp; English
                                                                                                                                                                                                                                                                                                                                                           240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВР.
                                                                                                                                                                                                                                                                                                                                                                                  GGAUGCUUCCAGGGCUC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95WO-US009080.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              134/c
AAT32134 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENP-) GEN-PROBE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1996-105849/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ryder TB, Kwoh TJ;
               WPI; 1996-105849/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9602557-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-FEB-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT32134;
                                                                                subjects.
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT32134/
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Gaps

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AAD19719;
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AAD19719/
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                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is an oligonucleotide complementary to an antisense oligonucleotide specific for the HIV target site, tat, which can be used for the detection of HIV, or for the treatment of HIV infection. The antisense oligonucleotide has an average EC(90) (mM) of 1500, which refers to the conc. of oligonucleotide required to achieve 90 % inhibition of HIV p24 core antigen prodn. (contg. phosphorothicate
                                                                                                                                                                                                                                                                                                                                                               corresponding to HIV sequences - used for the for inhibiting HIV propagation, partic. in infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                  Gaps
                                                                                                                                                         Oligonucleotide complementary to HIV tat targetting antisense oligo
                                                                                                                                                                            Human immunodeficiency virus, HIV; antisense oligonucleotide; tat; detection; treatment; infection; inhibition; p24; core antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 1; Length 21;
Pred. No. 3.2e+02;
0; Mismatches 2; Indels
3.2%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 3.2e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 6 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Simian herpesvirus B DNA primer B4.
                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 63; 90pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     240 GGCTGCTTCCCGGGCTC 256
                                   240 GGCTGCTTCCCGGGCTC 256
                                                                                                                                                                                               production; complementary; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.2%;
                                                                                                                                                                                                                                                                     95WO-US009080
                                                                                                                                                                                                                                                                                       94US-00277857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; 21
                                                      GGATGCTTCCAGGGCTC
                                                                                                  AAT32109 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 88.2
Matches 15; Conservative
       Best_Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                         (GENP-) GEN-PROBE INC.
                                                                                                                                                                                                                                                                                                                                                               Oligo:nucleotide(s)
detection of HIV or
                                                                                                                                                                                                                                                                                                                            Ryder TB, Kwoh TJ;
                                                                                                                                                                                                                                                                                                                                              WPI; 1996-105849/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV33173 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linkages only)
                                                                                                                                                                                                                                                                                       19-JUL-1994;
                                                                                                                                                                                                                                                                     14-JUL-1995;
                                                                                                                                                                                                                                  WO9602557-A1
                                                                                                                                       16-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-1998
                                                                                                                                                                                                                                                    01-FEB-1996
                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                   subjects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV33173;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                     AAT32109;
                                                     17
Query Match
                                                                                  RESULT 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 21
AAV33173
ID AAV3
XX
AC AAV3
XX
DT 06-N
XX
DE Simi
                                                                                           AAT32109,
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The invention provides the Simian herpesvirus B DNA (AAV33167) sequence coding for a gB glycoprotein (UL27; AAW70293) and a portion of an ICP 15. Kba protein (UL28; AAW70294). The invention uses these DNA and protein sequences as a basis for the development of differential diagnostic tests for the rapid identification of Simian herpesvirus B cases. Primer BV1 (AAV33169) and BV2 (AAV33169), along with the Simian herpesvirus B sequence specific PBs probe (AAV33179), were used in these diagnostic tests. Other primer sets used were the sense primers B3 (AAV33171) or B3' (AAV33172) and antisense primers B4 or B4' (AAV33174). Therefore, the virus can be detected by detecting the DNA sequence and knowledge of the amino acid sequence will help in the design of DNA production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monkey herpes B virus DNA - coding for gB glycoproteins and polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, Mammary Gland Cancer Specific Gene, MSG, cytostatic, vaccine; cancer, therapy, immune response, PCR primer; ss.
Simian herpesvirus B gB glycoprotein; UL27; ICP protein; UL28; differential diagnostic test; immunoassay; antibody; PCR; primer; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.2%; Score 13.8; DB 1; Length 21; 88.2%; Pred. No. 3.2e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 5 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Black
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eberle R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SWBI-) SOUTHWEST FOUND BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Col 7-8; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hilliard J, Scinicariello F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 TGACCGAGGCTGGGAC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-2000; 2000US-0192277P.
                                                                                                                               Synthetic.
Cercopithecine herpesvirus 1.
                                                                                                                                                                                                                                                                                                                                                          95US-00541878.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-2001; 2001WO-US009525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rcacceresecreseaec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-361791/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200172780-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                          10-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-OCT-2001.
                                                                                                                                                                                                                          US5767265-A.
                                                                                                                                                                                                                                                                                          16-JUN-1998.
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disease,

Wed Apr 21 12:58:21 2004

Hu P, Recipon H, Cafferkey R;

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Human polymorphic oligonucleotide U54701 fragment #14.
                                                                                                                                                                                    Sequence 21 BP; 3 A; 2 C; 9 G; 7 T; 0 U; 0 Other;
                                                                 Example 3; Page 77; 99pp; English
                                                                                                                                                                                                                         85 CAGTGGACATCACCACG 101
                                                                                                                                                                                                                                                                    AAH89013 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                           10-NOV-1999; 99US-0164596P.
                                                                                                                                                                                                                                                                                                                                                                                                               10-NOV-2000; 2000WO-US030766
                                                                                                                                                                                                                                   21 CACTAGACATCACCACG S
                                                                                                                                                                                                                                                                                            27-FEB-2002 (first entry)
                                                                                                                                                                                                             15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       (GLAX ) GLAXO GROUP LTD. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                          tissues and organisms
     (DIAD-) DIADEXUS INC.
                             WPI; 2001-616468/71.
                                                                                                                                                                                                                                                                                                                                                                                       WO200134840-A2
                                                       gland cancer.
                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-2001.
                  Salceda S,
                                                                                                                                                                                                                                                                                                                                                          Key
Variation
                                                                                                                                                                                                                                                                                AAH89013;
                                                                                                                                                                                                 Query Match
Best Local S
                                                                                                                                                                                                                                                        RESULT 216
                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                              AAH89013,
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Location/Qualifiers

replace(11,a)

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New use of irinotecan for preparation of compositions for treating cancer in subject having genome with variant allele comprising cytochrome P450, subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifedipine oxidase), polypeptide S (CYPAS) polymelociticatic (II). (I) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate
                                                                                                                                                                                                                               The present invention relates to human oligonucleotides comprising a single nucleotide polymorphic site (SNP: AAH88797-AAH89219). The present sequence is one such oligonucleotide. The oligonucleotides can be used in forensics, paternity testing, crecation of polymorphisms with phenotypic traits, genetic mapping of phenotypic traits and marker assisted breeding of animals and crop plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide cytostatic; PCR primer; ss.
                                                                                  New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly and also in forensics and paternity testing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 3.2%; Score 13.8; DB 1; Length 21; Local Similarity 88.2%; Pred. No. 3.2e+02; lose 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 2 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure, Page 32, 86pp, English.
ä
Thomas
                                                                                                                                                                                        Claim 68; Page 11; 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             75 GAGGGCCGCGCAGTGGA 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JUL-2002; 2002WO-EP008219.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17 dagececercaereda 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACF62203 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-OCT-2003 (first entry)
Patil N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ä
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                                            WPI; 2001-335945/35.
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Chen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-FEB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACF62203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            셤
                                                                                                                                                                                                                                                                                                            The present sequence is a PCR primer used for amplifying human mammary gland cancer specific gene (MSG) cDNA. MSG is useful for diagnosing, gland cancer specific gene (MSG) cDNA. MSG is useful for diagnosing, catecting, monitoring, staging, prognosticating, imaging and treating mammary gland cancer in a patient by determining the levels of MSG in cells, tissues or bodily fluids in a patient and comparing the determined levels of MSG with levels of MSG in cells, tissues or bodily fluids from a normal human control, where a change in determined levels of MSG in the patient versus normal control is associated with the presence of mammary control is used for identifying potential therapeutic agents for use in imaging and treating mammary gland cancer. MSG antibody conjugated to a cycloxic agent is useful for treating mammary gland cancer in a patient. MSG vaccine is useful for inducing an immune control managing and its protein and for treating mammary gland cancer in a patient. MSG and its protein and for treating mammary gland cancer in a patient. MSG and its protein are seful as diagnostic markers for mammary control and cancer and for diagnosis and treatment of disorders of cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                     New isolated polynucleotide, mammary gland cancer specific gene (MSG), useful for diagnosing, monitoring, staging, imaging and treating mammary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, single nucleotide polymorphic; SNP; forensic science;
paternity testing; phenotypic trait; genetic mapping; animal breeding;
plant breeding; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/gtandard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
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dosage and/or an appropriate derivative of (1). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of targ resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the exemplification of the present invention
   888688888888
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Sequence 21 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;

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Gaps
                                ;
3.2%; Score 13.8; DB 1; Length 21; 78.9%; Pred. No. 3.2e+02; tive 1; Mismatches 3; Indels
                                                              336 GACCAGGCCGGCTGCTCT 354
                                                                                           21 GTCCTGGGCCKGCTGCTGT 3
                  15; Conservative
   Query Match
Best Local S
                                  Matches
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Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide 5; cytostatic; PCR primer; ss.
                                       Cancer based on CYP3A5 related oligonucleotide SEQ ID NO:3.
         ACF62202 standard; DNA; 21 BP
                             (first entry)
                             08-OCT-2003
                                                                     Synthetic.
                   ACF62202;
RESULT 218
    ACF62202
```

23-JUL-2002; 2002WO-EP008219. 23-JUL-2001; 2001EP-00117608. 24-MAY-2002; 2002EP-00011710. WO2003013534-A2. 20-FEB-2003,

Heinrich G, Kerb R;

(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

WPI; 2003-268144/26.

New use of irinotecan for preparation of compositions for treating cancer in subject having genome with variant allele comprising cytochrome p450, subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.

Disclosure; Page 32; 86pp; English.

The present invention describes the use of irinotecan (1) or its derivative for the preparation of a pharmaceutical composition for treating colorectal, gastric, lung, ovarian or panceatic cancer, or malignant glional as aubject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifedipine oxidase), polypeptide (1779AS) polymotechide (11). (1) and (II) have cytostatic activity. The therapeutic applications of (1) is improved, since it is possible to individually treat a subject with an appropriate derivative of (1). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of treatment with substances (nonresponders), as well as the development of change resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62701 and ABM34312 to ABM3613 represent sequences used in the

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Gaps ö

3.2%; Score 13.8; DB 1; Length 21; 78.9%; Pred. No. 3.2e+02; Iv Mismatches 3; Indels

78.98;

Local Similarity 78.9 les 15; Conservative

Matches

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Query Match

336 GACCAGGGCCGGCTGCTCT 354

21 GTCCTGGGCCKGCTGCTGT 3

RESULT 220

Sequence 21 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;

Sequence 21 BP; 0 A; 6 C; 8 G; 6 T; 0 U; 1 Other;

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                                                                                                                                                                                                                                                                                                                                                    irinotecan; colorectal cancer; cervical cancer; gastric cancer; lung cancer; ovarian cancer; pancreatic cancer; malignant glioma; variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1
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     Length 21;
   3.2%; Score 13.8; DB 1; Length 2:
78.9%; Pred. No. 3.2e+02;
ive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                MRP1 based cancer related nucleic acid SEQ ID NO:4.
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                                                                            336 GACCAGGCCGGCTGCTCT 354
                                                                                                                1 Greeresecensers 19
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                        ADB20874 standard, DNA; 21
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3.27
Best Local Similarity 78.99
Matches 15; Conservative
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                                                                                                                                                                                                                                            ADB20874;
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ADB20873;

ADB20873

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The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTAL gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTALA gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancraatic cancer or malignant glioma. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.2%; Score 13.8; DB 1; Length 21; 78.9%; Pred. No. 3.2e+02; ive 1; Mismatches 3; Indels
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ovarian cancer; pancreatic cancer; malignant glioma; uridine diphosphate glycosyltransferasel member Al.
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24-MAY-2002; 2002EP-00011710.
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nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide.
                                                                                                                                                                                                                   irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;
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                                                                                                                                                                             MRP1 based cancer related nucleic acid SEQ ID NO:3.
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24-MAY-2002; 2002EP-00011710.
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Best Local Similarity
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New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising
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24-MAY-2002; 2002EP-00011710
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                                        WPI; 2003-289896/28.
                             Heinrich G, Kerb R;
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The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gateric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance I (MDR1) polymucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the
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lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; human; ds; Cyp3A5; MRP1; MDR1;
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                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 0 A; 6 C; 8 G; 6 T; 0 U; 1 Other;
multidrug resistance 1 polynucleotide.
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                                          Disclosure; Page 69; 130pp; English
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24-MAY-2002; 2002EP-00011710.
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ID ADB9
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                                                                                                                                                                                                     Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
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                                                                       (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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       23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
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The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant gliona in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant

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the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention.
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                                                                                                                                                       irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
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                                                                                                                    Human UGI1A1 variant allele sequence fragment SEQ ID NO:3.
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24-MAY-2002; 2002EP-00011710.
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  ADB92136 standard; DNA; 21
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Best Local Similarity 78.9
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kerb R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-342400/32.
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10-MAR-2003
20-APR-1993
                                                                                                                                                                                                                                              Homo sapiens
                                                                           04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ31318;
                                      ADB92136;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel use of irinotecan or its derivative for cervical, properation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidarug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; ds; human; UGT1A1; MRP1; TOP1.
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                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human UGT1A1 variant allele sequence fragment SEQ ID NO:4.
                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;
                                                                             BP; 6 A; 8 C; 6 G; 0 T; 0 U; 1 Other;
                                                                                                                Query Match 3.2%; Score 13.8; DB 1;
Best Local Similarity 78.9%; Pred. No. 3.2e+02;
Matches 15; Conservative 1; Mismatches 3;
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24-MAY-2002; 2002EP-00011710
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                                                                           Sequence 21
                                      invention.
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Gaps

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RESULT 226 ADB92136

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Feighner SD;

Elbrecht A,

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Comparative analysis of regions close to both the 3' and 5' ends of small subunit ribosomal RNA sequences with near identity in the eukaryotic kingdom identified two consensus sequences. RRIB 1 and BRIB 10, spanning the ssrRNA genes contained within the genomic DNA prepd. from a number of Eimeria species, to determine the degree of similarity between ssrRNA from different Eimeria species. The probe Common4RC represents a sequence common to all Bimeria species which may be used to identify Eimeria infection. See also AAQ31283-332. NOTE: As specifications EP-516395. BP-516395. Early specifications can be found indexed under EP-516395. However the claims section, sequences for all these specifications can be found indexed under EP-516395. However the claimed sequences of each specialication will be indexed under their own patent number, thus each separate patent will be represented. (Updated on 10-MAR-2003 to add missing OS field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Glutaminase, antiviral, virucide, anticancer; cancer therapy, HIV virus, gene therapy, Escherichia coli, primer; ss.
                                                                                                                                                                                                    Species-specific Eimeria tenella DNA probes - comprise divergent DNA sequences and are complementary to E. tenella small sub-unit ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pseudomonas glutaminase primer JR-1.
                                                                                                                                                                                                                                                                  Disclosure, Page 21; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               399 AAGGTCTTCTACGTGATCGA 418
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                                                                                                                        Chakraborty PR,
P-JuchelkaH;
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               92EP-00304781,
                                              91US-00707362
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(first entry)
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                                                                                        (MERI ) MERCK & CO INC.
                                                                                                                                                                        WPI; 1992-400736/49.
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                                                                                                                         Dashkevicz M,
                                                                                                                                          Liberator PA,
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               27-MAY-1992;
                                             29-MAY-1991;
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                                                            12-MAY-1992;
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12-JAN-1995
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Matches
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Oligo-nucleotide(s) able to inhibit multi:drug resistant phenotype - either by anti:sense or aptameric effects, useful for enhancing cytotoxic effects of chemotherapeutic agents on multi:drug resistant cancer cells.
                                                         Recombinant glutaminase derived from Pseudomonas 7A - expressed in E. coli to increase yield and avoid Pseudomonas endotoxins for antiviral and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/*tag= aakbone selected from: phosphorothioate;
/note= "Backbone selected from: phosphorister; morpholino
dithioate; methylphosphonate; phosphodiester; morpholino
backbone; polyamide backbone, and any combination of
these backbone types; the backbone may be modified to
incorporate a ribozyme structure, or a pendant group"
                                                                                                                                                Chromosomal DNA from Pseudomonas sp. 7A (ATCC 29598) was used to construct a genomic library in Escherichia coli LE392. Screening with mixed oligonucleotide probes was used to isolate a glutaminase encoding clone. This was sequenced using the primares given in AAQ68439-47. The gene can be used to manufacture recombinant glutaminase, free of Pseudomonas exotoxin, for use in e.g. HIV and cancer therapy. The gene may also be used in gene therapy protocols. (Updated on 25-WAR-2003 to
                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human multidrug resistance-1; MDR-1; inhibition; aptameric; human multidrug resistance-associated protein; antisense; cytotoxic;
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                                                                                                                                                                                                                                                                                                                         20;
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                                                                                                                                                                                                                                                                                                                         Length
Freeman
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                                                                                                                                                                                                                                                                                                                      Score 13.6; DB 1;
Pred. No. 3.2e+02;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;
Sethuraman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                      Disclosure; Fig 2B; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                    265 TGCACCTGGAGCAGGGGGGG
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                                                                                                                                                                                                                                                                                                                         3.2%;
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Roberts J, Macallister TW,
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nes 16; Conservative
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                             WPI; 1994-217891/26.
                                                                                          anticancer therapy
                                                                                                                                                                                                                                                               correct PN field.)
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misc_feature
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Matches
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The polypeptides of the invention may be of use in treating these
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vaccine, eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                        The present sequence represents a novel oligonucleotide OL(8E)MRP that specifically hybridises in a human cell with a complementary sequence of human multidrug resistance-associated protein (MRP) gene. Hybridisation causes inhibition of expression of the multidrug resistance phenotype by the cell, due to the oligonucleotide having an aptameric inhibitory effect as well as an antisense inhibitory effect. The oligonucleotide season and an antisense inhibitory effect. The oligonucleotide is administered to cancer patients to prevent development of the multidrug resistant phenotype. When co- administered with chemotherapeutic agents, the oligonucleotide is useful for potentiating elimination of multidrug resistant tummour cells from bone marrow or peripheral stem cell grafts.
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                                                                                                                                                                                                                                                          Also, the oligonucleotide can be used as an immunosuppressive agent
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                                                                                                                                                                                                                                                                                                                                         Score 13.6; DB 1; Length 20; Pred. No. 3.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                 4; Indels
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    Disclosure; Page 17; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                            28 AGGGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 AGGGCGGGATGATGATGGC 20
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97FR-00016034.
98US-0107077P.
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17-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs anchode purple that the genome of Chlamydia trachomatis. Antiennes and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases uch as conventional trachom, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal unetritis, energidymitis, cervicitis, salpingitis, perimpgatitis, batholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccine, eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
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                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                    20;
                                                                                                     4; Indels
                                              Length
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Seguence 20 BP; 3 A; 4 C; 6 G; 7 T; 0 U; 0 Other;
                                                 Score 13.6; DB 1;
Pred. No. 3.2e+02;
                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 1483; 1755pp; English.
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                                                                                                                                                            120 AAGTACGGCATGCTGGCCCG 139
                                                                                                                                                                                              20 AAATACGCCATGCTGACCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97FR-00015041.
97FR-00016034.
98US-0107077P.
                                                                                                                                                                                                                                                                                                                                                        멾.
                                                 Query Match 3.2%;
Best Local Similarity 80.0%;
                                                                                                                                                                                                                                                                                                                                                     AAZ01938 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                           16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chlamydia trachomatis.
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17-DEC-1997;
04-NOV-1998;
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amplification; hybridisation; organ transplant; gene typing; diagnosis;
ss.
                                                         WO200031295-A1
                                       Homo sapiens
                                                                                                07-OCT-1999;
                                                                                                                  26-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-SEP-2003
14-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUL-2000
                                                                            02-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lama glama.
                                                                                                                                                                                                                             diagnosis.
                                                                                                                                                         Moribe T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA73749;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 234
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                                                                                                                                                                                                                                                                                                                                                                                          AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX34584 - AAX35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae especially where the vector directs the expression of a neutralising
                                                                                                                                    Respiratory disease, pneumonia, bronchitis; heart disease, sarcoidosis, sinusitis; purulent otitis media; erythema nodosum, pharyngitis, vaccine, neutralising epitope; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human leukocyte antigen; HLA; class I allele type; probe; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human leukocyte antigen C allele DNA probe 361T368g SEQ ID NO:125.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                       Genome seguence of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                          Page 1724; Disclosure; 1912pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           389 CGGCGCCAAGAAGGTCTTCT 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CGTCACCAAGAAGTTCGTCT 20
         20 CTGCTTCCTTGGCACGCGA
                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA67067 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                   97FR-00014673
98US-0107078P
                                                        AAX95138 standard; DNA; 20
                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
                                                                                                                                                                                     Chlamydophila pneumoniae.
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epitope of C. pneumoniae
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                                                                                                                                                                                                                                                                                                                                     WPI; 1999-357842/30.
                                                                                                                                                                                                                                                                                               (GEST ) GENSET
                                                                                                                                                                                                                                                                   21-NOV-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-OCT-2000
                                                                                                                                                                                                                                               20-NOV-1998;
                                                                                                                                                                                                         WO9927105-A2
                                                                                               13-SEP-1999
                                                                                                                                                                                                                             03-JUN-1999
                                                                                                                                                                                                                                                                                                                 Griffais R;
                                                                                                                                                                           Synthetic.
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                                                                            AAX95138;
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                                     RESULT 232
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                                               AAX9513
                                                                   0x2x5x6x8
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The present invention describes a method for distinguishing a human belackocyte antigen (HLA) class I antigen or allele by a combination of polymerase chain reaction (PCR) using a primer pair whereby all HLA-A, -B or -C alleles can be amplified or using reverse hybridisation analysis comprising a DNA probe covalently bonded to microtire plate wells which are hybridisable specifically with the base sequence of at least one specific HLA-A, -B or -C allele. The method is applicable in gene typing, judying donor-recipient compatibility during organ transplant and correlation analysis for diagnosis of various diseases. The method is simple, rapid and accurate, with possibility of mechanisation and techniques. AAA666943 to AAA666977 represent oligonucleoride probes and PCR primers for use in the method of the present invention
                                                                                                                                                                                                                                                                                          Simple, rapid and accurate method for distinguishing HLA class I allele type with possibility of mechanization and automation, applicable in judging donor-recipient compatibility during organ transplant and disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression library; antibody; immunization; anchor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.2%; Score 13.6; DB 1; Length 20;
80.0%; Pred. No. 3.2e+02;
iive 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; & C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        187 CACATATCCACTGCTCGGTG 206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 8; Page 78; 83pp; Japanese.
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98JP-00335151.
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ID AAA73749 standard; DNA; 20
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                                                                           (SHIO ) SHIONOGI & CO LID
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Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                   Kaneshige T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
                                                                                                                                                                                                                          WPI; 2000-400097/34.
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framework; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200043507-A1.
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The invention relates to the human angiotensingen (AGT) gene, some mutant alleles of which cause a susceptibility to insulin-dependent diabetes semilitus (IDDM, type I diabetes). The AGT gene is located on chromosome 1442-43, a region linked to IDDM. The invention discloses genomic sequences comprising an alternative AGT gene (AAC91600-C291604) and a genomic sequence comprising an alternative AGT gene (AAC91600-CC (AAC91606). The invention also encompasses the specifically claimed human AGT mutant nucleic acid sequences AAC91667-C91684, and the mutant AGT alleles or gene products thereof which are related to detecting mutant AGT alleles or gene products thereof which are related to detecting mutant AGT alleles or gene products thereof which are related to IDDM, determining whether a person has, or is at risk of developing diabetes via detection of a polymorphism in the AGT gene, and methods of screening for drug candidates which may be useful in the treatment of diabetes resulting from an AGT mutation. Methods of preventing or treating diabetes are claimed which comprise the administration of a compound which agonises or antagonises wild-type or mutant AGT, which agonises a transgenic non-human animal, or cell line derived cerpensation, or which cleaves AGT proteins. In addition, the invention cencompasses a transgenic non-human animal, or cell line derived computators can be used to treat or prevent diabetes mellitus. AGT medulators can be used to treat or prevent diabetes mutant AGT gene createsing or frament used in an exemplification of the invention of alternative exon 1 PCR primer used in an exemplification of the invention
                        Novel angiotensinogen gene, mutant alleles of which causes susceptibility to insulin-dependent diabetes mellitus useful for diagnosis of predisposition to diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; mouse, SACI, carbohydrate; sweetener; ethanol; alcoholism; ss; obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas; blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy; protein replacement therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.2%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 3.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels
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                                                                                                                                         Example 2; Page 33; 83pp; English.
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28-JUL-2000; 2000US-0221419P.
10-NOV-2000; 2000US-0247443P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200183749-A2.
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AAS97449/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    synthetic or semi-synthetic nucleic acid sequences, not cloned from an municade source, where the nucleic acid sequences are derived from municades source, where the nucleic acid sequences are derived from municadenised immunoglobulins that are naturally devoid of light chains. The library is useful for the preparation of antibodies having binding specificity for a target antigen which avoids the need for a donor to have been previously immunized with the target antigen. The recombination of heavy and light chains is avoided, therefore preventing the formation of molecules that are non-functional. The number of hypervariable residues in the binding domain is reduced, allowing a more complete repertoire of possible binding variants to be obtained. The present sequence is a PCR primer targeted to anchor regions in liama antibodies. The primers (AAA73745 to AAA73745) amplified the framework regions P1, P2, F2c, F3 and F4. (Updated on 15-8EP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                   Expression library comprising nucleic acids not cloned from an immunized source, derived from immunoglobulins naturally devoid of light chains, use for producing antibodies specific for a target antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human angiotensinogen gene; ACT; insulin-dependent diabetes mellitus; type 1 diabetes; chromosome 1q42-43; single nucleotide polymorphism; IDDM; SNP; diagnosis; susceptibility; transgenic animal; drug screening; antidiabetic; gene therapy; alternative exon 1; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human angiotensinogen gene alternative exon 1 PCR primer, SEQ ID NO:17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to an expression library comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 3.2%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 3.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 29; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       269 CCTGGAGGGGGGCACCA 288
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                                                                                                                                                                     Van Der Logt CPE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAC91615 standard; DNA; 20 BP
99EP-00300351.
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                                                                                                                 (HIND-) HINDUSTAN LEVER LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcgrail M, Russell DL,
                                                       (UNIL ) UNILEVER PLC. (UNIL ) UNILEVER NV.
                                                                                                                                                                                                                                  WPI; 2000-482910/42.
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                                                                                                                                                                        Frenken LGJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1-MAY-1999;
19-JAN-1999;
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AAC91615;

RESULT AAC9161

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Gaps .. 0

× 검 Ś ij De Jong PJ, Chatterjee A, Tordoff MG; Bachmanov AA, Beauchamp GK, Ohmen JD, Reed DR, Ross D,

(MONE-) MONELL CHEM SENSES CENT

WPI; 2002-075162/10.

Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.

Claim 14; Page 75; 239pp; English.

The invention relates to an isolated polypeptide, comprising a variant form to f mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obseity, diabetes or alcoholism associated with SAC1 expression. The sequences are useful in screening for drugs and sweeteners. Recombinant cell lines and transgenic or repress function of SAC1. Predisposition to diabetes, obseity or alcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 gene, a sequence variation of the SAC1 lous may indicate a predisposition to diabetes, obseity and/or alcoholism and may provide a diagnostic mark. The polynuclectide can be detected in a biological sample by contacting the DNA with a probe to form a hybridisation complex which is then detected. The sequence represent cDNA encoding human and mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes

Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

Gaps ö Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; les 16; Conservative 0; Mismatches 4; Indels Query Match

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RESULT 237 ABL41764

ABL41764 standard; DNA; 20 BP. ABL41764;

29-MAY-2002 (first entry)

PCR primer used to amplify N-RAS proto-oncogene exon 2.

N-RAS; single base substitution; DNA mutation; PCR primer; ss.

Homo sapiens.

Synthetic.

US6346386-B1. 12-FEB-2002.

29-SEP-2000; 2000US-00677045

29-SEP-2000; 2000US-00677045

(ARUP-) ARUP INST.

Elenitoba-Johnson KSJ;

WPI; 2002-224990/28.

Determining mutation in DNA, comprises attaching guanine-cytosine-rich clamp to DNA, fluorescently labeling DNA and mixing it with denaturant,

heating to melt DNA and comparing melting temperatures of DNA and its wild type.

Example 3; Col 10; 16pp; English.

PCR primers ABL41762-64 were used to amplify exon 2 of the N-RAS protonocogene, in the course of the invention. The specification describes a method for determining whether a DNA sequence contains an alteration. The method comprises attaching a DNA sequent comprising one or more copies of the DNA sequence to a guanine-cytosine-rich clamp, fluorescently labeling the DNA sequence to a quantine-cytosine-rich clamp, fluorescently labeling the DNA sequence of the DNA sequent and a wild type alteration in the DNA sequence. The method is useful for determining whether a DNA sequence contains an alteration. The method is useful for determining whether a DNA sequence contains an alteration. The method is usitable for detecting a mutation as small as a single base substitution in a relatively large DNA fragment. As the disparity in melting temperatures is most evident in a lower melting domain of a DNA fragment, it is possible to distinguish single base substitutions within lower melting domain

Sequence 20 BP; 9 A; 3 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ó Query Match 3.2%; Score 13.6; DB 1; Length 20; Best Local Similarity 80.0%; Pred. No. 3.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels

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RESULT 238 ABQ74079

ABQ74079 standard; DNA; 20 BP.

ABQ74079;

(first entry) 11-0CT-2002 Microsatellite typing and sequencing D6S291 5' primer.

Homozygous stem cell; major histocompatibility complex; MHC; HLA; human leukocyte antigen; immunotype; genotype; microsatellite; probe; gent cell; nottopic; neuroprotective; antigarkimsonian; vulnerary; cytostatic; antiarteriosclerotic; antiinflammatory; immunosuppressive; antianaemic; antidiabetic; tranquilliser; respiratory; cardiant; trauma; muscular; ophthalmological; gene therapy; genetic disease; cancer; cystic fibrosis; muscular dystrophy; cardiac condition; burn; myopathy; neurodegenerative disease; Alzheimer's disease; Parkinson's disease; multiple sclerosis; pertrumar repair; reconstruction; blindness; disbetes; autoimmune disease; anaemia; PCR primer; ss.

Synthetic.

WO200257429-A2. 

25-JUL-2002.

32-JAN-2002; 2002WO-US000107.

02-JAN-2001; 2001US-025881P.

(STEM-) STEMRON INC.

Yan WL;

WPI; 2002-575456/61.

Producing homozygous stem cells having a target genotype and/or immunotype from non-fertilized post-meiosis I diploid germ cells, suitable for diagnostic, therapeutic and cosmetic transplant and

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The present invention describes a method for producing homozygous stem

(HS) cells having a target genotype and/or immunotype from non-fertilised

post-meiosis I diploid germ cells by mitotically activating the germ

cells to develop multiple blastocyst-like masses, each of which contains

an inner cell mass (ICM) that is homozygous for the target genotype

and/or immunotype. The methods of the present invention are useful for

the production of HS cells utilised for diagnosis, therapeutic and

cosmetic transplantation, cell replacement and/or gene therapy, and the

treatment of various genetic diseases (cystic fibrosis, muscular

disease, Parkinson's disease and multiple solerosis), traumatic injuries

(post-trauma repair and reconstruction, limb replacement, spinal cord

injuries and burns), cancer disorders of the epithalium (blindness,

myopathy, atherosolerosis), Crohn's disease, alabetes, autoimmune

diseases and anaemia. ABQ74028 to ABQ74115 represent PCR primers and

sequence specific oligonucleotide (SSO) probes which are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              exemplification of the present invention
                                                                                                               Disclosure, Fig 7; 75pp; English.
treatment of various disorders.
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Sequence 20 BP; 3 A; 6 C; 7 G; 4 T; 0 U; 0 Other;

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Length 20;
                     4; Indels
 Score 13.6; DB 1;
Pred. No. 3.2e+02;
                       0; Mismatches
                                           126 GGCATGCTGGCCCGCCTGGC 145
                                                               1 gecárrcagecarecerese 20
 3.2%;
                       16; Conservative
            Local Similarity
   Query Match
              Best Loca
Matches
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Human oligonucleotide sequence. ABZ88298 standard; DNA; 20 BP (first entry) 17-OCT-2003 ABZ88298; RESULT 239 

Human, antisense; lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive, cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

sapiens Omor

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

Katz E, Pabalan J, Sandrasagra A, Ka ,, Shahabuddin S; (EPIG-) EPIGENESIS PHARM INC. Li Y, Sar Tang L,

Aguilar D;

WPI; 2003-229219/22.

Nyce JW, Miller S,

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure, SEQ ID NO 3540; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intiation codon, coding region, 5' or 3' end genomic flanking regions, continuation codon, coding regions, or segions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an continual antial and a second active agent comprising an continual antial and a second active, hypotensive, has antiinflammatory, attiallargic, antiasthmatic, hypotensive, cuse in attisense gene therapy. The composition may have a cuse in attisense gene therapy. The composition may have a composition are prophylactic or therapeutic respiratory effect of an antisinal area prophylactic or therapeutic respiratory effect of an antisinal ammatory steroid in a subject, for reducing levels of continuing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine corpus professional antisense or condition, lung allergies, or treating bronchoconstruction, cung surfactant in a subject is tissue, or treating bronchoconstruction, lung allergies, or a respiratory disease or condition, or the sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cut the printed at the printed are the prophylaction and the printed are the printed and the p

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Sequence 20 BP; 5 A; 7 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö 3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; iive 0; Mismatches 4; Indels Query Match
Best Local Similarity 80.0'

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Gaps

BP. ABZ92729 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ92729; RESULT 240 ABZ92729 

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; dyfostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

24-APR-2001; 2001US-0286137P. 23-APR-2002; 2002WO-US013135.

(EPIG-) EPIGENESIS PHARM INC.

Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D; Tang L, Shahabuddin S; Nyce JW, L Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 7971; 872pp; English.

first active agent comprising an oligomortectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genee encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinon. A composition of the invention has antiinflammatory, antiallargic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antificlammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed of specification, but was obtained in electronic format directly from WIPO The invention relates to a novel pharmaceutical composition, which has a at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;

0; Gaps 3.2%; Score 13.6; DB 1; Length 20; 80.0%; Pred. No. 3.2e+02; Live 0; Mismatches 4; Indels 125 CGGCATGCTGGCCCGCCTGG 144 1 decchretrecchecres 20 16; Conservative Similarity Query Match Local Matches à

RESULT 241

AB298765

ABZ98765 standard; DNA; 20 

ABZ98765;

17-OCT-2003 (first entry)

Human tryptase b oligonucleotdie sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone, antiinflammatory, antiallergic, antiasthmatic; hypotenaive, immunosuppressive, cytostatic, gene therapy, antisense gene therapy, respiratory; lung, adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, K. Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 14007; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of compositions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory, antiallergic, antiasthmatic, hypotensis ye, immunosuppressive, and cytostatic activity. The composition may have a sum inflammatory antiallergic, antiasthmatic, hypotensis ye, immunosuppressive, and cytostatic activity. The composition may have a comparisory, integrately and cytostatic activity. The composition may have a compressive, and cytostatic activity. The composition may have a correcting a replicatory, lung or malignant disease or condition, as so for reducing sensitivity, to adenosite, reducing or depleting levels of antinflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine creeptor, producing bronchodilation, increasing bronchocomstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed of percification, but was obtained in electronic format directly from WIPO at first wino int/mh/mh/sibed and electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences 

Seguence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;

Gaps ., 3.2%; Score 13.6; DB 1; Length 20; 30.0%; Pred. No. 3.2e+02; Ive 0; Mismatches 4; Indels 80.08; 16; Conservative Local Similarity Query Match Matches

RESULT 242 ACC62132,

ACC62132 standard; DNA; 20

ACC62132;

20-JUN-2003 (first entry)

Human alipoprotein B antisense oligonucleotide SEQ ID NO: 21

alipoprotein B; ApoB; antilipaemic; antiarteriosclerotic; antidiabetic; anorectic; cardiovascular; gene therapy; lipid metabolism; cholesterol metabolism; atherosclerosis; hyperlipidaemia; diabetes; type 2 diabetes; obesity; atherosclerosis; cardiovascular disease; glucose; antisense oligonucleotide; ss.

Synthetic.

WO2003011887-A2.

13-FEB-2003.

30-JUL-2002; 2002WO-US024247

01-AUG-2001; 2001US-00920033. 30-APR-2002; 2002US-00135985. 15-MAY-2002; 2002US-00147196.

(ISIS-) ISIS PHARM INC.

Crooke RM, Graham MJ;

WPI; 2003-268105/26.

New antisense oligonucleotides for modulating apolipoprotein B, especially for preventing or treating atherosolerosis, hyperlipidemia or diabetes, or for modulating glucose, cholesterol, lipoprotein or triglyceride levels. 

Example 15; Page 96; 160pp; English.

(ISIS-) ISIS PHARM INC.

Watt AT;

Gaarde WA,

WPI; 2003-559091/52

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The invention relates to a novel compound that is 8-50 nuclectides in length that is targeted to a nucleic acid molecule encoding and specifically hybridises with and inhibits the expression of a nucleic acid molecule encoding ApoB; or which specifically hybridises with at least an 8-nuclectide portion of an cartive acid molecule encoding ApoB; or which cartive atte on a nucleic acid molecule encoding ApoB; or which ancertic, and cardiovascular activity. The compound of the invention has antilipaemic, antisense oilgonucleotide is useful for treating an animal having a disease or conditions associated with ApoB, e.g. a condition involving an abnormal lipid metabolism, a condition involving an abnormal metabolism, atherosclerosis, or a condition involving an abnormal metabolism, atherosclerosis or cardiovascular disease). Obesity, atherosclerosis or cardiovascular disease). Obesity, atherosclerosis or cardiovascular disease). The new compound or the antisense oligonucleotide is also useful for modulating glucose levels in a human or diabetic animal, or for modulating serum cholesterol levels, lipoprotein levels (specifically VuDL, HDL or LDL) or serum triglyceride levels, contenting or delaying the onset of a disease or condution associated with ApoB, or the onset of an increase in glucose levels in the animal or invantion. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "OTHER= phosphorothioate backbone, where 1-5 and 16-20 are 2' methoxyethyl nucleotides. All cytidines are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Libylif, Nov., instruction binding protein-related protein 2; IGFBP-rP2; insulin-like growth factor binding protein-related protein 2; IGFBP-rP2; IGFBP-8; Hos24; ecogenin; acute lymphoblastic leukaemia; gene therapy; hyperproliferative disorder; cancer; pulmonary fibrosis; renal fibrosis; scleroderma; atherosclerosis; cytostatic; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense; human, ss; connective tissue growth factor; CTGF; chromosome 6g23.1; ctgrofact; fibroblast inducible secreted protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human connective tissue growth factor antisense oligo DNA (SeqID 51)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB25658 standard; DNA; 20 BP
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention
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Gaps

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This invention relates to novel methods for modulating the expression of connective tissue growth factor (CTGF) by antisense oligonucleotides.

CTGF has been mapped to human chromosome region 623.1, and is also known as ctgrofact, fibroblast inducible secreted protein, fisp-12, NOV2, insulin-like growth factor binding protein-related protein 2, IGFBF-FP2, IGFBF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                          New antisense oligonucleotides for modulating connective tissue growth factor expression, particularly useful for treating cancers (e.g. breast or prostate cancer), pulmonary or renal fibrosis, scleroderma or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PKA regulatory subunit RII alpha inhibitory oligonucleotide ISIS102782.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.6; DB 1; Length 20; Pred. No. 3.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                            Example 15; Page 85; 139pp; English.
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Best Local Similarity 80.0
Matches 16, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-511923/48.
                                                                                                                                                                                                                                                 atherosclerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACD44753;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD44753
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The invention relates to antisense compounds targeted to nucleic acids encoding protein kinase A regulatory subunit RII alpha. The antisense compounds are useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophlaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human protein kinase A regulatory subunit RII alpha inhibitory oligonucleotide New antisense compounds, useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha, and for treating a disease or condition associated with expression of PKA regulatory subunit RII The present invention relates to proteins that have been found useful increasing the breakdown efficiency of protein-containing substances. present sequence represents a primer of the invention. Gaps Proteins for increasing breakdown efficiency of protein-containing Umitsuki G, Hatamoto O, Hara S, Masuda T, Sano M, Machida M; ö Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; les 16; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 2 A; 7 C; 5 G; 2 T; 0 U; 4 Other; Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other; (NODA ) NODA INST SCI RES. (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY. Disclosure; Page 61; 77pp; Japanese. Claim 14; Col 43-44; 35pp; English. CGCGTGCTGGCGGCGGACGA 337 crcareceececececes 20 protein breakdown; ss; primer. 20-AUG-2002; 2002WO-JP008376. 21-FEB-2002; 2002JP-00045090 ADB46018 standard; DNA; 20 Primer #1 of the invention. 04-DEC-2003 (first entry) WPI; 2003-697623/66 WO2003070954-A1 28-AUG-2003. Synthetic. ADB46018; Query Match alpha. Matches

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The invention relates to a method of identifying one or more biomarker genes for a type of cells among a group of (m) different cell types, from a multiplicity of genes whose expression levels in cells of the group are measured using nucleic acid arrays, to generate a plurality of measurements of expression levels for the m types of cells, by comparing rank order for the gene and (m-1) for each gene and detarmining a rank order for the gene among the multiplicity. The method is useful in identifying biomarkers using nucleic acid microarrays. The biomarkers of skin may be used in molecular diagnostic and pathology applications in normal and abnormal tissues and cell. The biomarker genes may also be used as molecular targets for therapeutics of a disorder or a disease in thumans. This sequence represents a QRT-PCR primer used in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying biomarker genes using nucleic acid microarrays, useful for molecular diagnostic and pathology applications, comprises comparing the Gibbs-likelihood ratios for each gene and determining a rank order for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                        ss; primer; biomarker gene; gene expression; nucleic acid array; molecular diagnostic method; molecular target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
3.2%; Score 13.6; DB 1; Length 20;
Best Local Similarity 80.0%; Pred. No. 3.2e+02;
Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 5 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 38; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              104 TGACCGCGACCGCAGCAAGT 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dooley TP, Curto BV, Davis RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TGACCCGACCTCAGAGAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE14433/C
ID ADE14433 standard; DNA; 20 BP
XX
AC ADE14433;
                BP.
                                                                                                                                                                                                                            COL6Al forward qRT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                         10-FEB-2003; 2003WO-US003673.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-FEB-2002; 2002US-0354519P
316 ACCGCGTGCTGGCGGCGG
                                                                                                                     ADC46898 standard; DNA; 20
                                                                                                                                                                                           18-DEC-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INTE-) INTEGRIDERM INC
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                                                                                                                                                                                                                                                                                                                                                    WO2003067217-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention.
                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the gene.
                                                                                                                                                           ADC46898;
                                                                                                      ADC46898
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Gaps

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Query Match
3.2%; Score 13.6; DB 1; Length 20;
Best Local Similarity 72.2%; Pred. No. 3.2e+02;
Matches 13; Conservative 3; Mismatches 2; Indel8

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The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (1) at least 5 ribose residues 5 (1) in a 2-C-allyl modification at position 4 of the ENA, (iii) at least ten 2-O-methyl modifications; and (iv) a 3-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. The PNA's can also be used to treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis Ribozyme therapy impactes on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The end of the ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly effected.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arthritic condition; graft tolerance; immune response; target; cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic nucleic acid molecules having a hammer-head motif - used for
the treatment of arthritis, induction of graft tolerance or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.4; DB 1; Length 15; 60.0%; Pred. No. 1.9e+02; tive 5; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human B7-1 hammerhead ribozyme target SEQ ID NO:1189.
                                                                                                                                                                                                                                                                                                                                  Stinchcomb DT, Jarvis T, Draper K, Gustofson J, Usman N, Wincott F, M Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 2 A; 3 C; 5 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 10; Page 166; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                             95US-00432874.
95US-00434509.
95US-0000951P.
95US-000974P.
95US-00512861.
                                                                                           94US-00354920.
94US-00363253.
94US-00363254.
95US-00390850.
95US-00426124.
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                                                              95WO-US015516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX64557 standard; RNA; 15
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                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-300653/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention
                                                                                                                                                                                                                                                                                                                                    Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                             23 - DEC - 1994
17 - PRB - 1995
20 - AAR - 1995
04 - MAX - 1995
07 - JUL - 1995
07 - JUL - 1995
07 - JUL - 1995
05 - OCT - 1995
WO9618736-A2
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                                20-JUN-1996.
                                                              22-NOV-1995;
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AAX64557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arthritic condition; graft tolerance; immune response; target; cleavage; hammerhead xibozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheunatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention describes a compound (I) 8-80 nucleobases in length targeted to a nucleic acid molecule encoding hydroxysteroid 11-beta dehydrogenase 1. inhibiting expression of hydroxysteroid 11-beta dehydrogenase 1. The methods and compositions of the present invention are useful for treating disorders associated with hydroxysteroid 11-beta dehydrogenase 1 expression, such as osteoporosis, depression and metabolic disorders like obesity, diabetes, atherosclerosis and hyperlipidaemia. This sequence represents an antisense oligonucleotide used to control the expression of human hydroxysteroid 11-beta
                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compounds useful for treating disorders associated with hydroxysteroid 11-beta dehydrogenase 1 expression, such as osteoporosis, depression and metabolic disorders like obesity, diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
                                                            osteopathic; antidepressant; anorectic; antidiabetic; antiarteriosclerotic; antilipemic; antisense-therapy; hydroxysteroid 11-beta dehydrogenase 1; osteoporosis; depression; metabolic disorder; obesity; HSD11B1; diabetes; atherosclerosis; hyperlipidaemia; antisense technology; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .Match 3.2%; Score 13.6; DB 1; Length 20; Local Similarity 80.0%; Pred. No. 3.2e+02; es 16; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 6 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                HSD11B1 antisense oligonucleotide seq id 35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; SEQ ID NO 35; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGACTTCCTCACTTTCCTGG 378
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                                                                                                                                                                                                                                                                                                     19-APR-2002; 2002US-00126355.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX64555 standard; RNA; 15
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29-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-852782/79.
                                                                                                                                                                                                     US2003198965-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            dehydrogenase
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                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                          Freier SM;
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Pavco P; Matulic-Adamic J;

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cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-11ke Growth Factor I receptor; IGF1; pityriaais; IGF binding protein, IGFPP-2; IGFBP); diffammation; psortiasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; sclaroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; between the condition of the retina; ser

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic;

IGF-I oligonucleotide #4549.

(first entry)

30-MAR-2001

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The present invention describes a novel enzymatic nucleic acid (ENA)

(ii) a 2'-C-allyl modification at position. The ENA; (iii) at least 5 ribose residues

(iii) a 2'-C-allyl modifications; and (iv) a 3'-end modification. The ENA's

cc in inhibit collagenase and stromelysin production in the synovial

cc an inhibit collagenase and stromelysin production in the synovial

con inhibit collagenase and stromelysin production in the synovial

con inhibit collagenase and stromelysin production of arthritis,

con inhibit collagenase and stromelysin production of arthritis,

con inhibit collagenase and stromelysin production of arthritis,

con inhibit collagenase and stromelysin production of a donor to induce tolerance

con a recipient to an alloantigen presenting cells of a donor to induce tolerance

controlly and inclease of contracting autoimmune disease, and for

controlly allergies and other inflammatory conditions. The ENA's can also

controlly allergies and other inflammatory conditions. The ENA's can also

controlly allergies and other inflammatory conditions. The ENA's can also

controlly allergies and other inflammatory conditions. The expession of

stromelysin without introducing the non-specific effects upon gene

controlling allergies. Ribozyme required to affect a therapeutic treatment

concentration of ribozyme required to affect a therapeutic treatment

controlling and examplification of the

controlling and example the controlling and example to a therapeutic treatment

controlling and example to affect a therapeutic treatment

controlling and example to affect a therapeutic treatment

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controlling and example to affect a therapeutic treatment

controlling and example and
hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; collagenase; stromelyžali, synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Enzymatic nucleic acid molecules having a hammer-head motif - used for
the treatment of arthritis, induction of graft tolerance or treatment or
auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stinchcomb DT, Jarvis T, Draper K, Pavco P;
Gustofson J, Usman N, Wincott F, Matulic-Adamic J;
Thompson JD, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 2 A; 4 C; 4 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 10; Page 166; 307pp; English.
                                                                                                                                                                                                                                                                                                                                   94US-00354920.
94US-00363253.
94US-003063554.
95US-00426124.
95US-00432874.
95US-00432874.
95US-000951P.
95US-000951P.
95US-000951P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                                 Homo sapiens.
                                                                                                                                                                                WO9618736-A2
                                                                                                                                                                                                                                                                                                                                                                                       23-DEC-1994;
17-FEB-1995;
20-APR-1995;
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                                                                                                                                                                                                                                  20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAY-1995
04-MAY-1995
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Ameliorating the effects of a disorder, e.g. psoriasis, by administering W (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 90; 201pp; English.

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Edmondson SR;

Wraight CJ, Werther GA,

WPI; 2001-041421/05.

(MURD-) MURDOCH CHILDRENS RES INST.

99US-0140345P.

21-NUL-1299;

21-JUN-2000; 2000WO-AU000693

WO200078341-A1.

28-DEC-2000.

Homo sapiens

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The present invention relates to a method for ameliorating the effects of antisense colacters. The method comprises contacting the skin with an antisense oligomuclectide, (for Insulin-like Growth Pactor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligomuclectide which can be used to design the antisense oligomuclectides of the present invention (see AAF45151 and AAF45153-6150muclectides of the present invention (see AAF45151 and AAF45153-616thyosis, pityriasis, thus, plants, serborrhoea, kelotds, keratosis, incoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, hyperproliferation of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%; Score 13.4; DB 1; Length 15; 93.3%; Pred. No. 1.9e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 3 A; 1 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             355 ACAGCGACTTCCTCA 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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Best Local
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Gaps

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1; Indels

3.1%; Score 13.4; DB 1; Length 15; 60.0%; Pred. No. 1.9e+02;

5; Mismatches

402 GICTICTACGIGATC 416 

Local Similarity 60.0

Query Match Best Loca Matches AAF53589/c ID AAF53589 standard; DNA; 15 XX

RESULT 250

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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                           Habener JF, Thomas MK;
                                                                                                     WPI; 2001-381492/40.
                                                                                                                                                                                                                                                                                                                                                                   WO200192524-A2.
                                        WO200141786-A1.
                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                       10-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                             36-DEC-2001
                                                   14-JUN-2001
                                                                                                                                                                                                                                                                                                      ABN07569;
                               Rattus sp.
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Matches
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 7561; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hanzel DK,
                                                                                                                                                                                                               30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
                                                                                                                                                                                                                                                                                                   30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
65-FEB-2001; 2001US-0266860P.
                                                                                                                                                30-JAN-2001; 2001WO-US000661
30-JAN-2001; 2001WO-US000662
30-JAN-2001; 2001WO-US000663
                                                                                                                                                                                                                                                                                   0-JAN-2001; 2001WO-US000667
                  25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                                                                                                                                                 (AEOM-) AEOMICA INC.
                                                                                                                         04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ઠે
The invention relates to a method of treating deficiency of insulin, that involves administering a hedgehog protein or nucleic acid encoding the hedgehog protein. The hedgehog proteins not be used in the method are selected from sonic hedgehog (Shh), indian hedgehog (Ihh) and desert hedgehog (Dhh). The method is useful for treating deficiency of insulin in a patient afflicted with diabetes, by stimulating insulin production in pancreatic beta-cells PBO). It is also used to treat deficiency of Insulin in patient, by stimulating IDX-1 production in PBC. The hedgehog protein is useful for modulating IDX-1 gene expression or its protein in PBC. This is used to treat deficiency of PBC in a patient, by stimulating neogenesis form beta-cell pancreatic ductal precursor cells. Inhibitors of the hedgehog proteins are useful for suppressing secretion of insulin in a patient affilicted with hyperinsulinemia. Sequences AAF84001-4002 represent PCR primers for ampliying the rat Dhh cDNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating deficiency of insulin, IDX-1 or pancreatic beta cells in a patient by, administering a hedgehog protein, nucleic acid encoding the protein or cells expressing the protein.
                                     Insulin, hedgehog protein, sonic hedgehog, Shh; indian hedgehog; Ihh; desert hedgehog, Dhh; diabetes; panoreatic beta-cell; PBC; IDX-1; neogenesis; hyperinsulinemia; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7561.
Rat desert hedgehog (Dhh) cDNA fragment amplifying reverse primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 3.1%; Score 13.4; DB 1; Length 16; Local Similarity 93.3%; Pred. No. 2.2e+02; les 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16 BP; 2 A; 4 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 29; 63pp; English.
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                                                                                                                                                                                                                                                                08-DEC-2000; 2000WO-US033575.
                                                                                                                                                                                                                                                                                                          99US-0170282P.
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                                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP.
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Chen W, Shannon ME;

Rank DR,

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids in samples, as amplification substrates and quantify hospide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specific blomclecule and/or amount apecifically of hGDMLP proteins, as specific blomclecule and/or amount apecifically of hGDMLP proteins, as specific blomclecule capture probes for surface-enhanced laser describing in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and with the expression of hGDMLP-1 sequence in patients and in the screening of the chopy selection with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a chopy sequence and in the expression of the present invention. N.B. The sequence data for this patent did not form part of the printed sequence data for this patent did not form art of the printed sequence data for this patent form form art of the printed sequence data for this patent did not form art of the printed sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.1%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.5e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  387 GACGCCCCAAGAAG 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 gacgegeccaagaag 16
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Best Local Similarity 93.3<sup>3</sup>
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABN79929;
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Ronaghi M, Ekstroem B,
                                                   WPI; 2002-393849/42.
                 WO200220837-A2.
                                                               incorporation.
                                                                                                                                                     20-NOV-2003
                                                                                                                                                                            Homo sapiens.
            Homo sapiens.
                                                                                                                                                                                 EP1281758-A2
                      14-MAR-2002.
                                                                                                                                                 ADA99492;
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                                                                                                                                       RESULT 25
ADA99492
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is concoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome concoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome concoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome concoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome concoded at chromosome concoded and modification of modification concoders. The modification concoders associated with decreased or increased expression or activity of MDZ3. Consed by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic consed by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic consed by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic consed by mDZ3, MDZ4, MDZ7, or MDZ12. The nucleic consed by mDZ3, wMZ4, MDZ7, or MDZ12. The nucleic consedul in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc fingar protein; MD23; MD24; MD27; MD212; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                          New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
3.1%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 2.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human MDZ3 scanning oligonucleotide SEQ ID 479.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; SEQ ID NO 481; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA99490 standard; DNA; 17 BP.
                                                                            30-JUL-2002; 2002EP-00016874
                                                                                                                                02-AUG-2001; 2001US-00922181
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                                                                                                                                                                                                                                      Shannon M, Gu Y, Nguyen C;
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                                                                                                                                                                                                                                                                                         WPI; 2003-423107/40.
                                                                                                                                                                                   (AEOM-) AEOMICA INC
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                         05-FEB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or more nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing two or more variable sites are typed, where three or more primer extension reactions are performed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid molecule(s). The methods are particularly suited for identifying microbals species or their subtypes, and in typing procedures e.g. typing of polymorphisms, tissue typing or in clinical applications. The sequence specific target region of genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ1; chromosome 7g22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15g26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                Human, single nucleotide polymorphism, nucleic acid typing, primer, tissue typing; sequencing, angiotensin converting enzyme, ACE, ss.
Human angiotensin converting enzyme SNP-fragment Eu6 primer A063FS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 1 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pourmand N;
                                                                                                                                                                                                                                                                                                                                                                                             (PYRO-) PYROSEQUENCING AB.
(STRD ) UNIV LELAND STANFORD JUNIOR.
(GARD/) GARDNER R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 47; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA99492 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                         10-SEP-2001; 2001WO-GB004042.
                                                                                                                                                                                                                                                                                                                                                08-SEP-2000; 2000GB-00022069.
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Gaps

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30-JUL-2002; 2002EP-00016874. 02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC

Gu Y, Nguyen C;

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Query Match 3.1%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.5e+02; Matches 14; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human MDZ3 scanning oligonucleotide SEQ ID 402.
                                                                                                                                                                                                                                                                                                             Example 8; SEQ ID NO 479; 103pp; English.
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                                                                                        WPI; 2003-423107/40.
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                                Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA99413;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               256
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ADA99413
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(first entry)

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ3 is concoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome corrections of manufacturing a medicament for treating or preventing a disorder sesociated with MDZ1, MDZ7, and MDZ12 sequences are useful in therapy, or imanufacturing a medicament for treating or preventing a disorder sesociated with MDZ12, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as protein are useful as therapeutic agents for gene therapy or as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.5e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 2 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human MDZ3 scanning oligonucleotide SEQ ID 480.
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                                                      Example 8; SEQ ID NO 402; 103pp; English.
MDZ4, MDZ7 or MDZ12, e.g. cancer.
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Best Local Similarity 93.3%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP1281758-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 6P21.3-22.2, MDZ7 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ1, and MDZ12 sequences are useful in therapy, or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene chreapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                                                                                     New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The mucleic acids and proteins are also useful for disgnosting or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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Sequence 17 BP; 4 A; 3 C; 7 G; 3 T; 0 U; 0 Other;

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Query Match 3.1%; Score 13.4; DB 1; Length 17; Best Local Similarity 93.3%; Pred. No. 2.5e+02; Matches 14; Conservative 0; Mismatches 1; Indels
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292 TGGTGAAGGACCTGA 306 2 recresassascrica 16 셤

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ADA99412 standard; DNA; 17 BP. ADA99412 

20-NOV-2003 (first entry) ADA99412;

Human MDZ3 scanning oligonucleotide SEQ ID 401.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

05-FEB-2003

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 401; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7922.1, MD24 is encoded at chromosome 1502.1.3-22.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15p11.2 or preventing at chromosome 15p2 is concided at chromosome 15p12.2 is encoded at chromosome 15p12.2 is and MD212 is encoded at chromosome 15p2.2 or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are

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                                                                                           Gaps
useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                        Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ra8; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
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                                                                  Length 17;
                                                                                          1; Indels
                                           Sequence 17 BP; 2 A; 7 C; 2 G; 6 T; 0 U; 0 Other;
                                                                 3.1%; Score 13.4; DB 1;
93.3%; Pred. No. 2.5e+02;
ive 0; Mismatches 1;
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                                                                                                                                                                                                                                                                     Human HER2 DNAzyme substrate #597.
                                                                             Local Similarity 93.3%; Pre
                                                                                                                                                                                               ABZ65140 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                               29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                          29-MAY-2002; 2002WO-US016840
                                                                                                                 363 Trecreating 377
                                                                                                                                     Trocreactareers 16
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                                                                                                                                                                                                                        ABZ65140;
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                                                                     Query Match
                                                                                                                                                                           RESULT 259
                                                                                            Matches
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule immension immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-HER2, K-Ras, H-Ras, and HIV acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, blown in AbzS9889 - AbzG2216, AbzG4544 - AbzG5531, AbzG6520 - AbzG6524, AbzG6530 - AbzG6585 represent substrate/target sequences for the human Match 3.1%; Score 13.4; DB 1; Length 17; Local Similarity 80.0%; Pred. No. 2.5e+02; es 12; Conservative 2; Mismatches 1; Indels Sequence 17 BP; 2 A; 5 C; 6 G; 0 T; 4 U; 0 Other; ribozymes of the invention Query Match

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nes 14; Conserv
                                                                                                                                                                                                           (BMLB-) BML INC.
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Iwasaki T;
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                                                                  Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                               Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                    Murine oligonucleotide associated with tumour supression, SEQ ID 1117.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 4 A; 5 C; 3 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Tuijnder M;
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2 ccacggugcagcugg 16
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                                                                                   ACC63870 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                 (MOLE-) MOLECULAR ENGINES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-333167/31
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tes 14; Conserv
                                                                                                                                                                                                                                           schizophrenia; ss.
                                                                                                                                                                                                                                                                                                 WO2003025176-A2.
                                                                                                                                                                                                                                                                        Mus musculus.
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ABL90998
ID ABL9
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AC ABL9
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DT 27-M
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The present invention describes a method for detecting lipid metabolism errors in patients using as indicators a set of 65 specific low density lipoprotein (LDI) receptor gene mutations. The method can be used in the diagnosis of an inherited predisposition to the development of diseases associated with hyperlipaemia, such as arteriosclerosis and ischaemic heart disease. ABL91141 encodes the LDL receptor given in ABB90525. ABL91142 to ABL91183 represent PCR primers used in the amplification of the receptor gene. ABL909090 to ABL91140 and ABB90445 to ABB90524 represents sequences used in the exemplification of the present invention
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LDL-R;
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antiarrhythmic: gene therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.1%; Score 13.4; DB 1; Length 19;
ilarity 82.4%; Pred. No. 3.1e+02;
Conservative 1; Mismatches 2; Indels
Hominidae; low density lipoprotein receptor; LDL receptor; detection; lipid metabolic error; hyperlipaemia; mutation; arteriosclerosis; ischaemic heart disease; ischaemia; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            376 TGGACCGCGACGACGGC 392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-00135020
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                                                                                                                                                                                                                                                                                                                                                                                                     17-JUL-2001; 2001WO-JP006153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JUL-2000; 2000JP-00218039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ90684 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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Wed Apr 21 12:58:21 2004

encoding minK protein and KVLQT1 protein involved channel formation useful for screening drugs, for

forms of genes

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Splawski

Sanguinetti MC,

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The invention relates to KVLQT1 and KCNE1 genes, associated with long QT (LQT) syndrome. It provides a mink protein comprising a mutation which substitutes the wild type amino acids with Leu, Asp, Leu, His, Trp and Al ao Thr ar residues 74,76,28,32,98 and 127 respectively. Screening KVLQT1 and KCNE1 is useful for identifying mutations for diagnosing and treating LQT. The ability to predict LQT enables physicians to prevent the diseases with medical therapy such as beta blocking agents and opts for better treatments. Sequences AAZ90675-Z90706 represent human KVLQT1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human long QT syndrome-associated KVLQT1 exon 5/intron 5 boundary.
                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other;
                                                                  in cardiac potassium channel formation use:
preventing and treating cardiac arrhythmia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KVLQT1; mutation; human; cardiac I(k8)
cardiac arrhythmia; electrocardiogram;
chromosome llpl5.5; intron; exon.
                                                                                                     Example 11; Page 69; 167pp; English.
                                                                                                                                                                                                                         intron/exon junction sequences
                                                                                                                                                                                                                                                                                                                                                                                                     AAZ98914 standard; DNA; 20
                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 93.35
                                WPI; 2000-195262/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200006199-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-FEB-2000
           Keating MT,
                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ98914;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     intron
                                                                                                                                                                                                                                                                                                                                                                             RESULT 263
AAZ98914/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   exon
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BP.

(first entry)

The invention relates to KVLQT1 nucleic acids which have a mutation compared to wild-type KVLQT1 (AAZ98901) The KVLQT1 gene encodes a protein of 676 amino acids which forms a cardiac I(ks) poteassium channel with the KCNE1 protein (AAY80563). The KVLQT1 gene contains 15 introns and encodes a protein containing 6 putative transmembrane segments and a pore forming region. The gene has been mapped to the chromosomal location 11p15.5. The sequences AAZ98905-28935 represent the intron-exon boundaries from the KLVLQT1 genomic sequence. Mutations in the KVLQT1 or KCNR1 genes result in cardiac arrhythmias observed as a prolonged QT curve in electrocardiograms (Long QT syndrome). The genes and proteins can be used for the diagnosis of subjects with long QT syndrome. They can also be used to screen for drugs which can be used for treating or preventing long QT syndrome. The KVLQT1 nucleic acids can be used for gene therapy, Human, ss; PARP; Poly (ADP-ribose) polymerase; antisense oligonucleotide; cytostatic; nootropic; neuroprotective; antinfilammatory; antidiabetic; immunosuppressant; hyperproliferative disorder; cancer; cellular injury; oxidative stress; neurological disorder; parkinsonism; apoptosis; meningitis-associated intracranial complication; ischaemia; probe; inflammatory disorder; authritis; diabetes. New isolated mutant KVLQT1 nucleic acids, useful for developing products for the diagnosis, prevention and treatment of long QT syndrome. Gaps \*tag= b mod\_base= OTHER note= "All cytidine residues are 5-methyl cytidine" ·, 3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels /\*tag= d
/\*tag= d
/mod\_base= OTHER
/not== "2' methoxyethyl nucleotides" /note= "2' -methoxyethyl nucleotides" KVLQTī peptides can be used for peptide therapy note= "Phosphorothicate backbone" Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other; Human PARP-3 antisense inhibitor ISIS #126076. Location/Qualifiers Example 11; Page 72; 178pp; English base= OTHER OTHER BP. 01-MAR-2001; 2001WO-US006572. 136 CCCGCCTGGCGTGG 150 "mod base= AAS45876 standard; DNA; 20 \*tag≈ c (first entry) 15 CCCACCTGGCGGTGG Conservative 16. .20 pou/ Local Similarity WO200164955-A1 modified base modified\_base modified base modified\_base Homo sapiens 07-SEP-2001, 14; 18-DEC-2001 AAS45876; Query Match Matches AAS45876 RESULT 요 ઠે .. 0

potassium channel; KCNEl; ss; Long QT syndrome; gene therapy;

Location/Qualifiers

sapiens

/number= 5 11. .20 /\*tag= b

'number=

๙

...10 /\*tag=

Connors TD;

Landes GM,

Curran ME,

Sanguinetti MC,

Keating MT, Sanguine Burn TC, Splawski I;

WPI; 2000-195199/17.

(UTAH ) UNIV UTAH RES FOUND. (GENZ ) GENZYME CORP.

L7-AUG-1998;

99WO-US010260 98US-0094477P

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Gaps

.. 0

3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels

136 CCCGCCTGGCGGTGG 150

15

Wed Apr 21 12:58:21 2004

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The invention relates to antisense oligonuclectides targeted to human PARP nucleic acid and inhibiting expression of human PARP. PARP (Poly And PARP) and inhibiting expression of human PARP. PARP (Poly decondensation, DNA replication, DNA repair, gene expression, malignant transformation, cellular differentiation and apoptosis. The antisense oligonuclectide inhibitors are useful for inhibiting the expression of PARP in human cells or tissues. They are also useful for treating a human with a disease associated with PARP especially hyperproliferative disorders (e.g. cancer), cellular injury resulting from oxidative stress, environgations and ischaemia), inflammatory and autoimmune disorders (e.g arkinsonism, meningitis-associated intracranial complications and ischaemia), inflammatory and autoimmune disorders (e.g arthritis) and diabetes. The present sequence is an antisense oligonucleotide of the invention
                                                                                                                                                                                                                                                                             Antisense compound useful for treating hyperproliferative, neurological, inflammatory and autoimmune disorders and diabetes inhibits human PARP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 91; 168pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14; Conservative
(ISIS-) ISIS PHARM INC.
                                                                                                  Popoff I, Cowsert LM;
                                                                                                                                                                                          WPI; 2001-602570/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; tive 0; Mismatches 1; Indels
                                                                            273 GAGCAGGCGCGCACC 287
                                                                                                          SAGCAGGGCTGCACC 15
                                                                                                              g
                                                                              8
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Human KVLQTI exon/intron boundary for exon #5
     AAC89924 standard; DNA; 20 BP
                   (first entry)
                   08-MAR-2001
             AAC89924;
   AAC89924
RESULT
```

Human; KVLQT1; antiarrhythmic; cardiant; gene therapy; cardiac potassium channel; Jervell and Lange-Nielsen Syndrome; JLN; chromosome 11p15.5; long QT syndrome; ss.

Homo sapiens

JS6150104-A.

98US-00135021 17-AUG-1998; 31-NOV-2000

97US-00874655. 98US-0094477P. 13-JUN-1997; 29-JUL-1998;

(UTAH ) UNIV UTAH RES FOUND.

Splawski I;

Keating MT,

WPI; 2001-060013/07.

DNA encoding for a mutant KVLQT1 which causes Jervell and Lange-Nielsen syndrome (JLN) when homozygous, useful for diagnosing long QT syndrome, or diagnosing or prognosing JLN.

Example 5; Col 45-46; 58pp; English

KVLQT1 is a cardiac potassium channel and mutations in the KVLQT1 gene cause Jervell and Lange-Nielsen Syndrome (JLN). KVLQT1 maps to chromosome lip15.5. The present invention relates to a mutant KVLQT1 coding sequence (see AAC89914). The mutant KVLQT1 coding sequence is useful in the presence diagnosis of long off syndrome and in screening humans for the presence of KVLQT1 gene variants which cause JLN syndrome. The present sequence is an exon/intron boundary of KVLQT1 XXXXXXXXXXXXXXXXX

Sequence 20 BP; 2 A; 8 C; 8 G; 2 T; 0 U; 0 Other;

ö Gaps .; 0 Length 20; 1; Indels Match
Local Similarity 93.3%; Pred. No. 3.5e+02;
hes 14; Conservative 0; Mismatches 1; Query Match

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ВЪ. AAI69777 standard; DNA; 20 RESULT 266 AA16977

16S/23SrRNA spacer region PCR primer #3. 13-DEC-2001 (first entry) AA169777; 

Bacterium detection; 168/23SrRNA spacer region; PCR primer; ss.

; 0

Gaps

; 0

Pseudomonas putida. JP2001190279-A.

13-JAN-2000; 2000JP-00004160. 17-JUL-2001.

(MITO ) MITSUBISHI JUKOGYO KK 13-JAN-2000; 2000JP-00004160.

Detection method of Pseudomonas bacteria. WPI; 2001-605311/69.

Claim 9; Page 8; 11pp; Japanese.

The present invention relates to a method for the detection of the 163/23SrRNA spacer region of Pseudomonas putida (see AAI69774). The method can be used to detect Pseudomonas bacteria. The present sequence is a PCR primer which was used in an example from the present invention

Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;

ö 3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels Best Local Similarity 93.3%; Matches 14; Conservative Query Match

ö

Gaps

3 CCAGGAGTGAACTG 17 Ŋ CCAGCAGTGAAACTG 13

à

AAL40401 standard; DNA; 20 BP. RESULT 267 AAL40401 ID AAL4 XX AAL4

AAL40401

environmental monitoring; food industry; feed industry;

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Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoletic disorder; cancer; neurological; Alzheimer's disease;
                                                                                                                                                                                                                                                                  An antisense oligonucleotide of 8 to 50 nucleotides in length that inhibits caspase 6, is useful for treating Rieger's syndrome.
                                                                                                                                                                                                                                                                                                  Claim 3; Page 92; 141pp; English.
                                                                                       apoptotic; mouse; murine; ds.
                                                                                                                                                                   03-OCT-2001; 2001WO-US030871
                                                                                                                                                                                       04-OCT-2000; 2000US-00679299
                                                                                                                                                                                                                              Brown-Driver VL, Zhang H,
         19-SEP-2002 (first entry)
                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                 WPI; 2002-471315/50
                                                                                                                             WO200229066-A1.
                                                                                                           Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a deoxy gap
                                                                                                                                                 11-APR-2002
                             Mouse
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Watt AT;

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Favis R, Kliman R;

Gerry NP,

Sarany F, Zirvi M, WPI; 2002-034366/04

(CORR ) CORNELL RES FOUND INC. 14-APR-2000; 2000US-0197271P.

04-APR-2001; 2001WO-US010958

WO200179548-A2 25-OCT-2001

caspase 6 antisense inhibition related oligo SEQ ID No 120.

Synthetic

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The present invention describes a method (M1) for designing capture oligomucleotide probes (I) for use on a support to which complementary oligomucleotide probes (II) will hypridise with little mismatch, where coligomucleotide probes (II) will hypridise with little mismatch, where coligomucleotide probes (II) will hypridise with little mismatch, where coligomucleotide albeit is the method is useful to detecting infectious diseases caused by bacterial infectious agents consistent infectious agents consistent infectious agents consistent infectious agents consistent in the management of the present of the method is useful for detecting genets cirus, selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus consistent of method is also useful for detecting genetic diseases such the method is also useful for detecting genes or genes concer involving oncogenes, tumour suppressor genes, or genes betecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, muman papillomavirus types 16 and 18 and 11 ver cancers. The confident is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning celectron microscope and infrared microscope) the support at the food confident is also used for environmental and croscope) the support at the food confident in the support at the confident of particular sites and identifying if ligation of the oligomucleotide sequences. ABI82074 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antianflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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3.1%; Score 13.4; DB 1; Length 20;
Best Local Similarity 93.3%; Pred. No. 3.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 6 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Fig 29; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 175 ACGAGTCCAAGGCAC 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ91337 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16 Accadrccaagccac 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 269
ABZ91337/C
셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 invention specifically hybridises to and inhibits expression of caspase 6 therappeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or axaxia telangiecrasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, various types of cancer neurological conditions such as Alzheimer's disease and other deregulated apoptoric pathological conditions. This polynucleotide sequence represents a mouse caspase 6 oligonucleotide relating to the invention. NOTE: This phosphorothioate oligonucleotide sequence has 2' MOE wings and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCAl; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic;
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Gaps
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Gaps

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3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; tive 0; Mismatches 1; Indels

122 GTACGCCATGCTGGC 136

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14; Conservative

Matches

Similarity

Query Match Local Capture oligonucleptide Zip ID#1370 oligo #9.

(first entry)

16-FEB-2002

ABI94283;

ABI94283 standard; DNA; 20 BP

RESULT 268 ABI94283/c

Seguence 20 BP; 3 A; 5 C; 7 G; 5 T; 0 U; 0 Other;

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20-FEB-2001; 2001US-00789529
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus norvegicus.
Oryctolagus cuniculus.
                                                                                                                                                                                                                                                                            WPI; 2003-029937/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer, transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Canis familiaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JS2002132290-A1.
                                          WO200277228-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus.
      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sus scrofa.
Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fugu ripes.
                                                                                03-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABX75395;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  g
    The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in atrisense gene therapy. The composition may have a cuse in atrisense gene therapy. The composition may have a cuse in atrisense gratery in a subject, for reducing or depleting levels of for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of creceptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, a tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Once: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; Artemis gene; DNA repair factor; metallo beta-lactamase; RS-SCID; chromosome 10; severe combined immunodeficiency; SCID1; cancer; PCR;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                       Li Y, Sandrasagra A, Katz B, Pabalan J, Aguilar D;
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.1%; Score 13.4; DB 1; Length 20; 33.3%; Pred. No. 3.5e+02; lve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer used to amplify Human Artemis gene exon 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 10 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 6579; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.3%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                    23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               282 GGCACCAAGCIGGIG 296
                                                                                                                                                                                                                               24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                                   EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV72389 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCACCAGGCTGGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14; Conservative
                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                            WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-2003
                                                                      Homo sapiens.
                                                                                                                                                  31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ubiquinone
                                                                                                                                                                                                                                                                                                                               Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABV72389;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                            Nyce JW,
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RESULT 270

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Best Loca Matches

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PCR primers ABV72389-ABV72416 were used to amplify exons of the human Artemis gene. This gene encodes a V(D)J recombination and/or DNA repair factor that belongs to the metallo beta-Lactamass enperfamily, and whose mutations give rise to the human RS-SCID condition. The gene is localised to chromosome 10. The Artemis gene or its nucleic acid is useful for diagnosing or treating severe combined immunodeficiencies (SCIDs) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CNS; conserved non-coding region; 88; cytokine; interleukin 4; IL-4; interleukin 5; IL-5; interleukin 13; IL-13; chromosome 5q31; LCR; PCR; locus control region; interleukin gene cluster; transcription factor; human; mouse; dog; rat; bovine; pig; rabbit; fruitfly; puffer fish;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid molecule of the Artemis gene, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 93.3%; Pred. No. 3.5e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                    (INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                              Fischer A;
                                                                                                                                                                                                                                                                                                                                                                                              diagnosing or treating SCID or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 62; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABX75395 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Forward PCR primer for CNS-6.
                                                              22-MAR-2001; 2001WO-IB000546.
22-MAR-2001; 2001WO-IB000546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50 CCACTCAGAGGAGTC 64
                                                                                                                                                                                                                  De Villartay J, Moshous D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003 (first entry)
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The invention relates to an isolated nucleic acid molecule comprising a length of about 100 nucleotides of less, which has a sequence at least about 100 nucleotides of less, which has a sequence (CNS)-1 sequence (a locus control region (CRS) element in interleukin gene (CNS)-1 sequence (a locus control region (CRS) element in interleukin gene (CNS)-1 coluster region of chromosome 5431 containing interleukin (IL) 4, ILS and (CNS-16 or mouse CNS-16 or their complements. Also included are: (1) an expression cassette comprising a CNS-1 sequence operably linked to a promoter which controls transcription of a heterologous coding sequence (CNS-16 or mouse CNS-1 sequence) (3) an expression cassette consisting a cesentially of an IL-4 gene, an IL-13 gene, and a CNS-1 sequence flanked between two recombination site sequences; (3) an expression cassette comprising an IL-4 gene and an IL-13 gene, and lacking a CNS-1 sequence; (5) a T cell comprising one of the expression cassettes; (6) a non-human transgenic caninal where a CNS-1 sequence; (5) a T cell; animal comprising one of the expression cassettes or the T-cell; and (7) a non-human transgenic animal where a CNS-1 sequence is deleted from its chromosome. The T cell is useful for identifying a compound that compound on binding of the transcription factor to a CNS-1 sequence of the CNS-1 sequence of the CNS-1 sequence of the CNS-1 sequence of the compound on binding of the transcription factor of a compound. The nucleic acid is useful for modulating expression cassettes is compound. The nucleic acid is useful for modulate functions of CNS sequence of the CNS-1 sequence of the CNS-1 sequence of the compound of the modulate functions of CNS sequence.

Compound: The nucleic acid is useful for modulate functions of CNS sequence of the CNS-1 sequence and that a diagnostic tool to screen patients having consecuted to cytokine gene expression. Expression cassettes with and without CNS are identifying compounds that modulate and the compound of a serial expression cassettes and 
                                                                                                                                                                                                                                                                                                                                                             Novel isolated nucleic acids which are locus control region elements in interleukin gene cluster region of chromosome, referred as conserved non-coding sequences, useful for modulating expression of cytokine genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, ARTEMIS protein, V(D)J recombination, DNA repair; gene therapy, severe combined immunodeficiency; SCID; cancer; exon 1, PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     degenerate PCR primer used to isolate a CNS sequence from a variety of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              various therapeutic modalities. The present sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.4; DB 1; Length 20; 82.4%; Pred. No. 3.5e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Artemis exon 1 amplifying PCR primer, Ex1F1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 1 C; 7 G; 5 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 20; 48pp; English.
                                                                                                                                                                                             Loots GG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              360 GACTTCCTCACTTTCCT 376
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AAD47533 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18 GACATCCTCACTNTNCT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-FEB-2003 (first entry)
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                                                                                                                                                                                   Frazer KA, Rubin EM,
                                                                                                                                                                                                                                                                               WPI; 2003-165733/16.
(FRAZ/) FRAZER K A.
(RUBI/) RUBIN E M.
(LOOT/) LOOTS G G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD47533;
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The invention relates to an Artemis nucleic acid coding for a protein involved in V(D)J recombination and/or DNA repair. Sequences of the invention are useful for treating severe combined immunodeficiencies (SCID) or cancer. They are also useful for diagnositing a patient, including a prenatal diagnosis with SCID, a predisposition to cancer, an immune deficiency or a carriage of a mutation increasing the risk of progeny to have such a disease. Peptides of the invention are used for preparing antibodies. The invention is useful in gene therapy. The present sequence is a PCR primer used to amplify human Artemis exon 1 DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HLA; polymerase chain reaction; inflammatory arthropathy; susceptibility;
arthritis; arthritis related diseases; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                        New ARTEMIS nucleic acid coding for a protein involved in V(D)J recombination and/or DNA repair, useful for treating and diagnosing severe combined immunodeficiencies (SCID) or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.1%; Score 13.4; DB 1; Length 20; 93.3%; Pred. No. 3.5e+02; ive 0; Mismatches 1; Indels '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                         (INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                     De Villartay J, Moshous D, Fischer A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BRBI-) BRITISH BIO-TECHNOLOGY LTD.
                                                                                                                                                                                                                                                                                                     Example 1; Page 66; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human leukocyte antigen probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ24900 standard; DNA; 18 BP.
                                                                                                                             22-MAR-2001; 2001WO-IB000546.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90GB-00024005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91WO-GB001935.
                                                                                                21-MAR-2002; 2002WO-IB001737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50 CCACTCAGAGGAGTC 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 CCAATCAGAGGAGTC 6
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 93.3
hes 14; Conservative
                                                                                                                                                                                                               WPI; 2003-018886/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1992-183691/22
                                        WO200277026-A2.
              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9207956-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-NOV-1990;
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19-NOV-1992
                                                                   03-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAY-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ24900;
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AAQ24900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Sequence 18 BP; 2 A; 7 C; 5 G; 4 T; 0 U; 0 Other;

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AAQ56855,
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셤 ઠે žS ਨੇ 셤 The sequence is that of a probe which hybridises to one of the human leukcyte antigen (HLA) sequences in the primer extension products (or strands) produced during PCR amplification of the HLA class I alleles. It is specific for the sequence encoding amino acids 67-71 (CKAKA) of the alpha 1 domain of the HLA-B27 group and is thus specific only for this group. It can be used in the detection and/or identification of an HLA sequence that may be indicative of a patients succeptibility to inflammatory arthropathy such as arthritis and arthritis related diseases. Such diseases include reactive arthritis, rheumatoid arthritis, Reiter's syndrome, uveitis, viral arthritis, psoriatic arthropathy, gouty arthritis, septic arthritis, erythema nodosum, Henoch-Schloelein purpura and esp. ankylosing spondylitis. See also AAQ24895-Q24902. (Updated on 25-WAR-2003 to correct PV field.) ö Sets of PCR primers (see AAQ56835-Q56857) are used as probes to detect Norwalk-related viruses, e.g. SRSV/KY/89, HuCV Sapporo, HuCV Houston and primate calcivirus. Detection of viral RNA is by RT-PCR. (Updated on 25-MAR-2003 to correct PN field.) DNA from Norwalk and related viruses - used for preparing prods. for use in diagnostic assays, detection and vaccines for Norwalk and related amplification of nucleic acids using buffer soln. and chelating agent or detecting HLA class I alleles for determining susceptibility to Norwalk virus; HuCV; Sapporo; pathogen; acute gastroenteritis; food poisoning; seafood contamination; diagnostic assay; PCR primer; human calcivirus; small round virus; polymerase chain reaction; ss. Gaps . Query Match
3.1%; Score 13.2; DB 1; Length 18;
Best Local Similarity 83.3%; Pred. No. 3e+02;
Matches 15; Conservative 0; Mismatches 3; Indels PCR primer P-74 for detection of Norwalk-related virus. Sequence 18 BP; 6 A; 6 C; 5 G; 1 T; 0 U; 0 Other; Graham DY; Claim 49; Page 104; 156pp; English. Disclosure; Page 13; 52pp; English Jiang X, BAYU ) BAYLOR COLLEGE MEDICINE 173 CTACGAGTCCAAGGCACA 190 78 AAQ56855 standard; DNA; 18 BP. 93WO-US008447. 92US-00941365. CTGCAAGGCCAAGGCACA (revised)
(first entry) Matson DO, Estes MK, WPI; 1994-101125/12. for detectin arthritis etc. WO9405700-A2 07-SEP-1993; 07-SEP-1992; 25-MAR-2003 05-OCT-1994 17-MAR-1994 Synthetic. AAQ56855; viruses.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cleavage of resistant DNA sites with restriction enzymes - using activator comprising recognition site and cleavage-permitting flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                               Gaps
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Score 13.2; DB 1; Length 18; Pred. No. 3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA cleavage; restriction endonuclease; NaeI; activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 0 A; 5 C; 10 G; 3 T; 0 U; 0 Other;
                            Pred. No. 3e+02;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Col 21; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                               'n
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                                                                                               271 TGGAGCAGGCGCCACCA 288
                                                                                                                                                                                                                                                                                                                                                                                                               NaeI substrate oligonucleotide
                                                                                                                                                                                                                                                        ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93US-00128369.
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       Query Match 3.1%;
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                                                                          18 redadcadeccadecrea
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                                                                                                                                                                                                                                                           AAQ87132 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
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Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conrad MJ, Topal MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  recognition site; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1995-199738/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-DEC-1990;
                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
06-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-MAY-1995.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                      AAQ87132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 276
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                                                                                                                                                                                                                                     AAQ87132
                                                                                                                                                                                                                  RESULT
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(first entry)

93JP-00260984 93JP-00260984

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Distinguishing different human herpes virus strains
                                                                                                                                                                                                                        Seguence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                   Cytomegalovirus detection oligonucleotide #3.
                                                                                                                                                                                                                                                                                                                                                      Human herpesvirus group B primer #1
                                                                                                                                                                     Claim 1; Page 9; 10pp; Japanese.
                                                                                                                                                                                                                                                                216 AACTCGGTGGCGGCCAAA 233
                                                                                                                                                                                                                                                                                                                                                                                 sandwich hybridisation; ss
                                                                                                                                                                                                                                                                                                                AAT01523 standard; DNA; 18
                                                                                                                                                                                                                                       Query Match
Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-370480/48
                                                                                                                                   WPI; 1995-196320/26.
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                                                                                                                      (TOYM ) TOYOBO KK.
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                                                                                                                                                                                                                                                                                                                                                                                                            JP07250699-A
                                                                                            19-OCT-1993;
                                                                                                        19-0CT-1993;
                                                                 JP07111893-A
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                                                                                                                                                                                                                                                                                                                                                                                                                          03-OCT-1995.
     12-JAN-1996
                                                                              02-MAY-1995.
                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                    Synthetic.
                                                                                                                                                                                                                                                                               18
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                                                                                                                                                                                                                                                                                                  RESULT 277
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ВЪ.

(first entry)

94JP-00041101 94JP-00041101

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Primers and probes AAT01515-40 and AAT16978-87 are used in a novel method for the specific detection of human herpes viruses (HHV) in which at least two types of HHV nucleic acids are pre-amplified by at least 4 primers, followed by a separate detection step using specific detection probes. The primers and probes are synthesised based on the sequences of at least 8 HHV strains selected from HSV1, HSV2, VZV, EBV, CMV, HHV-6A, HHV-6B and HHV-7. They are split into 3 groups: A, B or C. Similarly the probes are split into 3 groups: A', B' and C'. The probes are specific in that they will only detect the amplification prods. from that virus by sandwich hybridisation. This primer is derived from Epstein-Barr virus by and cytomegalovirus (CMV) sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         An oligonucleotide for the amplification and the specific detection of Epstein-Barr virus (EBV) and cytomegalovirus (CMV) - useful for detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q876296-Q876303 are PCR primers used in a new method for the amplification and specific detection of Epstein-Barr virus (EBV) and cytomegalovirus (CMV). The oligonucleotides are useful for the detcting the EBV and CMV genes from a culture supernatant of herpes virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
amplification with at least 4 primers and hybridisation to specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer; oligonucleotide; Epstein-Barr virus; cytomegalovirus; CMV; amplification; detection; herpes; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Epstein-Barr virus (EBV) and cytomegalovirus (CMV) PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 3.1%; Score 13.2; DB 1; Length 18; Best Local Similarity 83.3%; Pred. No. 3e+02; Matches 15; Conservative 0; Mismatches 3; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 3e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 4 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                 Claim 3; Page 10; 14pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      216 AACTCGGTGGCGGCCAAA 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   216 AACTCGGTGGCGGCCAAA 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 6; 7pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and in diagnostic procedures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93JP-00273615.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-211626/28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ87296;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 278
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ID AAO8
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                                                                                                             Cytomegalovirus; hybridisation assay; radioisotope; fluorescent compound; enzyme; linker arm; biotin; RNA polymerase promoter; immobilisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The oligonucleotides AAQ92471-86 can be used for the detection of cytomegaloviruses in a hybridisation assay. The oligonucleotides may be modified by labelling with radioisotopes, fluorescent compounds, enzymes, nucleotides with linker arms, biotin or the promoter sequence for an RNA polymerase. The oligonucleotides may be optionally immobilised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer, PCR, amplification, probe, human, herpes virus, cytomegalovirus, herpes simplex virus, varicella zoster virus, Epstein-Barr virus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligo:nucleotide(s) for detection of cytomegalovirus - can be with labels, useful in hybridisation assays, opt. immobilised.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%; Score 13.2; DB 1; Length 18; 83.3%; Pred. No. 3e+02; ive 0; Mismatches 3; Indels
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AAX09121-X1026B are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a
                                                                                                                                                                                                                                  forms (AAT41710-13) of the murine major histocompability complex
                                                                                                                                                                                                                                      Mutated forms (AAT41710-13) of the murine major histocompability complex interferon-stimulated response element (MMC IRSE) binding sequence (AAT41709), along with other 'competitor' DNAs (AA73174-16), were used in gel shift assays designed to determine whether mouse lymphocytenespecific interferon regulatory factor (LSIRF) (see also AAR99426) is a DNA binding protein. Mutant misRE mutant mt4 (AAT41713) competed well with wild-type MHC ISRE for binding to LSIRF protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphism, biallelic; human; forensic; paternity testing; disease; detection; phenotypic typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; ss.
                                                                                                                                          New genes for murine lymphocyte specific interferon regulatory factor used for modulation of lymphocyte activation and proliferation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated nucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity testing or phenotypic typing for disease.
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
3.1%; Score 13.2; DB 1; Length 18;
Best Local Similarity 83.3%; Pred. No. 3e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human biallelic polymorphic marker downstream primer #393.
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 7 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                     Richardson CD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WHED ) WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 16; Page 197; 310pp; English.
                                                                                                                                                                                                   Example 4; Page 41; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 GCCAGGAGTGAAACTGCG 19
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96US-00611280
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                                                                       Grossman A,
                                   (AMGE-) AMGEN CANADA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lander ES, Wang D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-286974/25.
                                                                                                          WPI; 1996-477128/47.
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Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9820165-A2
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 03-APR-1996;
                                                                       Matsuyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAY-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX10087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ88011-30 are oligonucleotides used for the detection of Epstein-Barr virus. There is no cross reaction with other type of herpes viruses using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligo:nucleotide(s) for detection of Epstein-Barr virus - have no cross reactivity with other herpes viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lymphocyte specific interferon regulatory factor; LSIRF; IRF-3; probe; major histocompatibility complex; MHC; ISRE; interferon-stimulated response element; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                         Oligonucleotide probe 10 for detection of Epstein-Barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.1%; Score 13.2; DB 1; Length 18; larity 83.3%; Pred. No. 3e+02; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse MHC ISRE binding sequence mutant mt4.
                                                                                                                                                                                                                                                                probe; detection; Epstein-Barr virus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 9; 10pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               216 AACTCGGTGGCGGCCAAA 233
                                                                                                              AAQ88020 standard; DNA; 18 BP
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 18 ACCTTGGTGGTGGCCAAA
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                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             these probes
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                                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                                                                                                      AAQ88020;
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Matches
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Gaps

Page 137

cc method for determining polymorphic forms in an individual for use in e.g. forensics, paternity testing or for phenotypic typing for diseases such cas agammaglobulinemia, diabetes insipidus, Leschi-Nyham syndrome, muscular dystrophy Wiskott-Aldrich syndrome, Fabry's disease, familial cypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrand's disease, tuberous sclerosis, hereditary haemorrhagic telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria, cyptem, infection by pathogenic microorganisms, and characteristics such as longevity, appearance (e.g. baldness, obsesity), strength, speed, endurance, fertility, and susceptibility or receptivity to particular drugs or therapeutic treatments. The isolated polymorphic nucleic acid segments can also be used to produce medicaments for the treatment or prophylaxis of such diseases 8 $\pm$ 

Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 18; Pred. No. 3e+02; ); Mismatches 3; Indels ., 344 CCGCCTCCTCCACCGA 361 1 ccecrccreracaca 18 3.1%; Query Match
Best Local Similarity 83.38
Marches 15; Conservative 셤 8

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AAX80491 standard; DNA; 18 BP. AAX80491

AAX80491;

(first entry) 26-AUG-1999 Human secreted protein yc2\_1 probe.

Human, secreted protein, immunostimulator, nutrition, cytokine, cell proliferation, differentiation, immune stimulating, vaccine, suppression, haematopoiesis regulation, tissue growth, activin, inhibin, chemotactic, chemokinetic, haemostatic, thrombolytic, anti-inflammatory; cadherin; tumour invasion suppressor; tumour inhibition; gene therapy; probe; hybridisation; ss 

Homo sapiens. Synthetic

WO9932614-A1

01-JUL-1999.

98WO-US027140. 18-DEC-1998; 97US-0068379P. 98US-00212843. 20-DEC-1997; 16-DEC-1998;

GEMY ) GENETICS INST INC.

, Evans C; Wong GG, Clark HF; Collins-Racie LA, Steininger RJ, W Lavallie ER, Agostino MJ, Mccoy JM, Treacy M, Jacobs K, Merberg D, Fechtel K;

WPI; 1999-395405/33.

New polynucleotides encoding secreted human proteins potentially useful as, e.g. immunostimulators.

Disclosure; Page 96; 99pp; English.

The present invention describes human secreted proteins obtained from human fetal brain, fetal Kidney or adult blood cDNA libraries. The present sequence represents a probe for a human secreted protein. The human secreted proteins, and polynucleotides encoding them, are predicted human secreted proteins, and polynucleotides encoding them, are predicted

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to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, immune stimulating (e.g. as vaccines) or suppressing activity, haematopoiesis regulating activity, tissue growth haemacotatic, chemokinetic activity, haemostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity. The polynucleotides are also stated to be
                                                                                                                                                                                                                                                                                                                                                                                                                                                       G-alpha-13; human; inhibitor; cancer; antisense compound; therapy; ss.
                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense compound inhibiting expression of human G-alpha-13
                                                                                                                                                                                        Score 13.2; DB 1; Length 18; Pred. No. 3e+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                              Human G-alpha-13 antisense inhibitor ISIS# 20742.
                                                                                                                                                                 Sequence 18 BP; 2 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                   30 GGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                                                             1 Grcregascarcriesc 18
                                                                                                                                                                                                                                                                                                                                                 ВP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-00205860.
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                                                                                                                                                                                             Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative (
                                                                                                                                                                                                                                                                                                                                                   AAZ31793 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                       24-JAN-2000 (first entry)
                                                                                                                                           useful for gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                             AAZ31793;
                                                                                                                                                                                                                                                                                                                       RESULT 283
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ö This sequence represents an antisense inhibitor of the invention, and inhibits the expression of the human G-alpha-13 protein. The antisense compounds of the invention are of 8 to 30 nucleobases in length, that inhibits the expression of the human G-alpha-13. The antisense compound is useful for treating an animal, particularly humans, having or being prone to a disease or condition associated with the expression of G-alpha-13, such as cancer Gaps ö Score 13.2; DB 1; Length 18; Pred. No. 3e+02; 0; Mismatches 3; Indels Sequence 18 BP; 4 A; 6 C; 8 G; 0 T; 0 U; 0 Other; Claim 11; Col 38; 38pp; English. 103 CTGACCGCGACCGCAGCA 120 1 cedacceccacecades 18 3.1%; Query Match
Best Local Similarity 83.33

Page 138

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cerebral ischemic neuronal damage; traummatic brain injury; peripheral neuropathy; Alzheimer's disease; Huntington's disease; Parkinson-Plus syndrome; progressive Supramuclear Palsy; Olivopontocerebellar atrophy; pshy-brager Syndrome; damanian parkinsonism dementia complex; amyotrophic lateral sclerosis; memory impairment; neuronal disorder; neuropathy; ischemic stroke; acute brain injury; neuropathy; acute spinal cord injury; nervous system tumour; multiple sclerosis; eve disorder; PCR primer; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated polypeptides, used for treating e.g. neurodegenerative disease or disorder, neuronal damage or neuronal disorder of the peripheral nervous system, the medulla or the spinal cord.
                                                                                      PCR primer NBNint.sense for neublastin neurotrophic factor cDNA.
                                                                                                          Neurotrophic factor; neublastin; neurodegenerative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 33; Page 32; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Johansen TE, Blom N, Hansen C;
                                                                                                                                                                                                                                                                                                                                                         98US-0092229P.
98DK-00001048.
98US-0097774P.
98DK-00001265.
                   AAZ60571 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                        99WO-DK000384.
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                                                                 05-MAY-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-171013/15.
                                                                                                                                                                                                                                                                            WO200001815-A2.
                                                                                                                                                                                                                                                                                                                        35-JUL-1999;
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19-AUG-1998;
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                                           AAZ60571;
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RESULT 284
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per primers AAZ66571-72 were used to amplify cDNA encoding a neurotrophic factor designated neublastin. Neublastin is a member of the glial cell line-derived neurotrophic factor sub-class of the transforming growth factor-beta superfamily of neurotrophic factors. Neublastin exhibits high affinity for the GFR-alpha3-RET receptor complex. The polypeptides can be used for treating a neurodegenerative disease or disorder, certebral ischemic neuronal damage, traumatic brain injury, peripheral neuropathy. Alzheimer's disease, Huntington's disease, Parkinson's disease, Parkinson also plus syndromes, progressive Supranuclear Palsy, Olivopontocerebellar etrophy, Shy-Drager Syndrome, Guamanian parkinsonism dementia complex, amyotrophic lateral solerosis, memory impairment, or a neuronal disorder of the peripheral nervous system, the medulla or the spinal cord. They can also be used for treating various neuropathies. They can also be used for treating various neuropathies. They can also be used for treating various neuropathies. They can also be used for treating ischemic stroke, acute brain injury, acute spinal cord injury, nervous system tumours, multiple sclerosis, exposure to neurotoxins, metabolic diseases such as diabetes or renal dysquincions in his events. and damage caused by infectious agents, or various disorders in the eye

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Sequence 18 BP; 3 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
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3.1%; Score 13.2; DB 1; Length 18;

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Sequences AAZ49796-Z59835 represent antisense oligonuclectides targetted to the human Smad3 gene, which inhibit its expression. The antisense oligonuclectides were designed to target different regions of the human condainment of the condainment of the smad state. The smad protein are a family of the human condains which are involved in TGF-bets superfamily signal transduction. To proteins which are involved in TGF-bets superfamily of cytosolic on ligand binding, TGF-bets superfamily proteins (such as bone con ligand binding, TGF-bets superfamily proteins (such as bone con phosphorylate Smad proteins which then home or heterodimerise and translocate to the nucleus to activin and TGF-bets themselves) con translocate to the nucleus to activite target gene transcription. Smad3 carrials transcription factors, the pathway-restricted Smads, which are consulated by TGF-bets and activitae transcriptical Smad4 (USc013787-A, AAY69622), the complex being able to activate TGF-bets inducible transcription. The oligonucleotides of the invention are useful conditions associated with smad4 conditions approached by the are tumour formation, inflammation and certain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense inhibition of the human Smad3 gene, useful for diagnosing, preventing and treating conditions associated with Smad3 expression e.g.
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                          Gaps
                                                                                                                                                                                                                                                                            Human Smad3 phosphorothioate antisense oligonucleotide, SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                          Smad3; MADH3; hWAD3; JV15-2; TGF-beta signalling pathway;
transcription factor; expression inhibition; antisense therapy;
tumour formation; inflammation; antisense; ss.
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                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 1 A; 4 C; 12 G; 1 T; 0 U; 0 Other;
          Pred. No. 3e+02;
); Mismatches
83.3%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 11; Col 38; 31pp; English.
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                                                              45 GGCCACCACTCAGAGGAG 62
                                                                                         1 GGCCACCGCTCCGACGAG 18
                                                                                                                                                                                ВР
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83.3%;
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                                                                                                                                                                                AAZS9797 standard; DNA; 18
                                                                                                                                                                                                                                                 (first entry)
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Best Local Similarity 83.3
Matches 15, Conservative
                            15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert LM;
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            Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                  AAZ59797;
                                                                                                                                               RESULT 285
                                                                                                                                                                 AAZ5979
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The invention relates to novel human secreted proteins, the nucleic acids encoding them. The protein may exhibit cytokine, cell proliferation or cell differentiation activity or may induce production of other cytokines in certain cell populations and may exhibit immune stimulating or immune suppressing activity, which is useful for the treatment of various immune deficiencies and disorders e.g. severe combined immunodeficiency (SCID), autoimmune disorders e.g. multiple sclerosis, systemic lupus crythematosus, rheumatoid arthritis, autoimmune pulmonary inflammation. The proteins are also useful in the treatment of diseases and disorders including tissue, skin and organ transplantation and in graft-versue-host including tissue, skin and organ transplantation and in graft-versue-host cliseases (GVBD), in the induction of tumour immunity, myeloid or lymphoid cell deficiencies, wound healing and tissue repair, in the treatment of burns, inclsions and ulcers, as well as in treatment of pariodontal disease, osteoporosis or osteoarthritis, mediated by inflammatory processes, diseases, Huntington's disease, amylotrophic lateral central nervous system vessel e.g. stroke, sepsis, inflammatory bowel disease, ulcers, bone regeneration. The protein, having activin- or inhibin-related activities is useful as a contraceptive based on the ability of inhibins male mammals. The proteins and nucleic acids are also spermatogenesis in male mammals. The proteins and nucleic acids are also spermatogenesis in male mammals. The proteins and nucleic acids are also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Secreted human proteins, useful as vaccine for treating various diseases such as autoimmune disorders (e.g. multiple sclerosis), and nervous
                                                                                                                                                                                                                                                                                                                                   numani, section discontantia antimization, immunically immunically interaction in contropic; neuroprotective, antiarthritic; antimicrobial, vulnerary, cytostatic; antidiabetic; virucide, antiinfertility; anticonvulsant; vasotropic; antiparkinsonian; immunecimulant; dermatological; probe; antithemmatic; antitumor; antitucer; osteopathic; tranquiliser; cerebroprotective; cytokine; cell proliferation; cell differentiation; immune deficiency; sCID; tumour; autoimmune disorder; multiple sclerosis; rheumatoid arthritis; autoimmune disorder; multiple sclerosis; neumatoid arthritis; paridontal disease; osteoporosis; os
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Wong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mccoy JM, Lavallie E, Collins-Racie LA, 1
Agostino MJ, Steininger RJ, Spaulding V,
Merberg D;
                                                                                                                                                                                                                                                                                                                  protein; ss; antiinflammatory;
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                                                                                                                                                                                                                                                    Human secreted protein yc2_1 probe.
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                                                         AAS59326 standard; DNA; 18 BP.
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04-DEC-2000; 2000US-00729674.
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                                                                                                                                                                                       16-JAN-2002 (first entry)
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Fechtel K,
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Treacy M,
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RESULT 286
AAS59326
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probe used to detect the nucleic acids of the invention and where an N residue is present at position 2 this is a biotinylated phosphoroamidite residue
                                                                                                                                                                                                                                                                                                                primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                SCR; sequence characterised amplified regions; beef; cow; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.1%; Score 13.2; DB 1; Length 18; 33.3%; Pred. No. 3e+02; ve 0; Mismatches 3; Indels
                                                                          Score 13.2; DB 1; Length 18;
Pred. No. 3e+02;
); Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                         SCR primer 1 for distinguishing between beef types.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 3 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                       G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scr primer for distinguishing korean beef meat.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hong YH, Jung IJ, Kim HB, Kim HS, Kim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 5; 6pp; Korean.
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                                                                                                                                30 GGCTGGGACGAAGATGGC 47
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Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                              Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative
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                                                       BP; 2 A; 3 C; 8
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                                                                                                                                                                                                                   ABL53448 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                             KR2001017747-A.
                                                                                                                                                                                                                                                                                                                                                       Unidentified.
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                                                        Sequence 18
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neuroprotective; haemostatic; thrombolytic; anti-inflammatory; phosphoarmidite; ss.
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(CLAR/) CLARK H.
(FECH/) FECHTEL K.
                                                                                   US2001039335-A1.
                                                                                                                                                                                                                                                                                                        30-JAN-1998;
18-FEB-1998;
23-NOV-1998;
                                                                                                                                               34-DEC-2000;
                                                                                                                   08-NOV-2001
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07-JAN-1998
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13-JAN-1998
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                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jacobs K,
                                                                                                                                                                                                                                                                                                                                                                                                                                               (EVAN/)
(MERB/)
(TREA/)
(AGOS/)
(STEI/)
(SPAU/)
                                                                                                                                                                                                                                                                                                                                                                                                  (MCCO/)
(LAVA/)
(COLL/)
                                                                                                                                                                                                                                                                                                                                                                                     (JACO/)
        The invention relates to a truncated neublastin polypeptide comprising an amino acid terminus that lacks one or more amino-terminal amino acids of a mature neublastin polypeptide. The polypeptides and mucleic acids are useful for treating neurodegenerative disorders such as ischemic neuronal damage, traumatic brain injury, peripheral neuropathy, neuropathic pain, Alzheimer's disease, Huntington's disease, Parkinson's disease, amyotrophic lateral sclerosis, memory impairment, diabetes, renal diseases, or glaucoma by moderating metabolism, growth, differentiation or survival of a nerve or neuronal cell. This polynucleotide sequence is a neublastin PCR primer of the invention
                                    Nootropic; neuroprotective; antiparkinsonian; anticonvulsant; analgesic; tranquiliser; antidiabetic; ophthalmological; neurodegenerative disorder; neublastin; ischemic neuronal damage; traumatic brain injury; diabetes; peripheral neuropathy; neuropatho; pain, Alzheimer's disease; glaucoma; Huntington's disease; Parkinson's disease; amyotrophic lateral sclerosis; memory impairment; renal disease; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                   New truncated neublastin polypeptides lacking one or more amino-terminal amino acids of a mature neublastin polypeptide useful for treating neurodegenerative disorders, e.g. peripheral neuropathy, neuropathic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.1%; Score 13.2; DB 1; Length 18; Best Local Similarity 83.3%; Pred. No. 3e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 3 A; 8 C; 6 G; 1 T; 0 U; 0 Other;
          Neublastin DNA related PCR primer SEQ ID No
                                                                                                                                                                                                                                                                                                                                                          Rossomando A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 44; 138pp; English.
                                                                                                                                                                                                                                             12-MAR-2002; 2002WO-EP002691
                                                                                                                                                                                                                                                                             12-MAR-2001; 2001US-00804615
                                                                                                                                                                                                                                                                                                                                                          Sah DWY, Johansen TE,
                                                                                                                                                                                                                                                                                                         (BIOJ ) BIOGEN INC. (NSGE-) NS GENE AS.
                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-713515/77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pain, brain injury
                                                                                                                                                                                   WO200272826-A2.
                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                 .9-SEP-2002.
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97US-0126425P. 97US-0067454P. 97US-0068379P. 98US-0070346P. 98US-0070643P.

2000US-00729674

98US-0070755P. 98US-0071304P. 98US-0072134P.

98US-0073095P 98US-0075038P 98US-00197886 2000US-00539330

MCCOY J M. LAVALLIE E R. COLLINS-RACIE I

JACOBS

EVANS C. MERBERG D. TREACY M. AGOSTINO M J. STEININGER R J

SPAULDING V.

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The invention relates to isolated polymucleotides (ABA90876-ABA90968 and ABA9080) and encoded proceins (ABB55698-ABB55800), especially controlled the account of the accoun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or systemic inflammatory response syndrome, ischaemia-reperfusion injury
                                                                                                                                                                                                                                                                                                                                                                    New secreted proteins and encoding polynuclectides, useful in gene therapies, particularly for preventing or treating autoimmune disorders, cancer, graft-versus-host disease, wound, osteoporosis, stroke or inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 329; 349pp; English
Lavallie ER,
1, Agostino MJ,
Fechtel K;
                                                              D, Treacy M,
Clark H, Fe
                                                                                                                                                                                                                                                                  WPI; 2002-040725/05.
                                                                            Merberg D,
Wong GG,
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Human; clone bd306-7; clone yb8-1; ATCC number 98599; gene therapy; immune disorder; bacterial infection; fungal infection; cancer; tumour; autoimmune disorder; systemic lupus erythematosus; wound; ulcer; inhibin; osteoporosis; osteoarthritis; nervous system disorder; neuropathy; Alzheimer's disease; Parkinson's disease; Huntington's disease; activin; Alzheimer's disease; Parkinson's disease; Huntington's disease; activin; ischaemia reperfusion injury; inflammatory bowel disease; chemotactic; crohn's disease; cytostatic; anti-inflammatory; immunomodulator;

Biotinylated oligonucleotide SEQ ID NO 213.

(first entry)

14-FEB-2002

ABA90995;

Collins-Racie LA, Evans C; Steininger RJ, Spaulding

Mccoy JM,

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Gaps . 0

62

GGCCACCACTCAGAGGAG

GGCCACCGCTCCGACGAG 18

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ABA90995 standard; DNA; 18

289

ABA9099 RESULT

Endoplasmic reticulum stress competence control element SEQ ID NO:11.

(first entry)

23-DEC-1999

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Gaps

AAZ25638;

踞

AAZ25638 standard; DNA; 19

RESULT 29 AAZ25638/

Endoplasmic reticulum; ER; stress competence; control element; inhibition; growth; apoptosis; cancer; autoimmune disease;

cystic fibrosis; ds.

JP11243959-A.

Gallus sp.

14-SEP-1999.

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The present sequence represents a specifically claimed oligonucleotide PCR primer. The oligonucleotide can be used for polymerase chain reaction (PCR) amplification of DNA, specifically regions of specific genes that are conserved among mammalian species, i.e. pairs of oligonucleotides from the present specification represent universal mammalian sequence-tagged site (UM-STS) primers. The primers are used to develop genomic maps, to isolate clones from libraries, to make cross-species comparisons and to develop additional genetic markers. UM-STS allow genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New oligonucleotide primers amplifying gene regions conserved among mammals - useful for developing genomic maps, isolating clones and making cross-species comparisons.
endotoxin lethality, arthritis, inflammatory bowel disease or Crohn's disease; or tumours or cancers, pemphigus vulgaris or pemphigus foliaceus. The present sequence is that of a biotinylated oligonucleotide with a phosphoaramidite residue, useful to the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; polymerase chain reaction; amplification; UM-STS; universal mammalian sequence tagged site; genomic map; clone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Interleukin 2 receptor PCR primer for universal mammalian STS's.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02;
                                                                                                                              DB 1; Length 18;
                                                                                                                                                                  3; Indels
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                                                                                           Sequence 18 BP; 2 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                          Query Match
3.1%; Score 13.2; DB 1
Best Local Similarity 83.3%; Pred. No. 3e+02;
Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yuzbasiyan-Gurkan V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 10; 26pp; English.
                                                                                                                                                                                                   30 GGCTGGGACGAAGATGGC 47
                                                                                                                                                                                                                                     78
                                                                                                                                                                                                                                                                                                                                 AAV01209 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-0012061P.
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(UNMS ) UNIV MICHIGAN STATE.
                                                                                                                                                                                                                                       1 Grciggacgargrigge
                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Brewer GJ, Venta PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-435083/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                        23-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9731012-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-AUG-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                    AAV01209;
                                                                                                                                                                                                                                                                                                 RESULT 290
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New control element for stress competence of endoplasmic reticulum - useful for inhibition of growth and induction of apoptosis in cancer

98JP-00052453 98JP-00052453

04-MAR-1998; 04-MAR-1998; (HSPK-) HSP KENKYUSHO KK.

WPI; 1999-603708/52.

Example 1; Fig 3; 25pp; Japanese.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention specifically claims an element shown by: (A) a 19 bp base sequence, CCAATNNNNN NNNNCCACG (ERSB); or (B) a modified base sequence having replaced 1.3 bases with the other base(s), which induces transcription with stress on endoplasmic reticulum used for stress competence of endoplasmic reticulum. Also described are: (1) a DNA having transcription inducing activity with stress on endoplasmic reticulum containing the above mentioned element, optionally further containing a promoter DNA, and (2) a vector containing the element can be used for the inhibition of growth and induction DNA. The element can be used for the inhibition of growth and induction dispasses and cystic fibrosis by inhibition of symptoms of canter cells, and improvement of symptoms of canter can be used for the inhibition of symptoms of autoimmune diseases and cystic fibrosis by inhibition of autoantibody formation.

AAZ25632 to AAZ25657 represent elements used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Endoplasmic reticulum; stress; ER; transcription factor; transcription; regulatory element; ERSE; bZIP; chaperone; treatment; prophylaxis; cancer; arteriosclerosis; ischaemia; wound healing; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.1%; Score 13.2; DB 1; Length 19; Similarity 83.3%; Pred. No. 3.46+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 4 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GRP94 promoter ERSE3-like sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         150 GAGGCCGGCTTCGACTGG 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВЪ.
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AAA28576 standard; DNA; 19
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
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29-AUG-2000
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Best Local S:
Matches 15
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Gapa

3; Indels

0; Mismatches

350 GCTCTACAGCGACTTCCT 367

Query Match Best Local Similarity 83.3 Matches 15; Conservative

18

GCTCTACAGAGAGGTCCT

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chicken; gene expression; GRP;

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An endoplasmic reticulum stress transcription factor (bZIP) capable of regulating transcription inducing activity exhibited by an element (ERSE) can be used in a method for controlling expression of an endoplasmic reticulum chaperone. The method comprises expression bZIP. The method can be used for expression of a foreign protein by positively regulating expression of an endoplasmic reticulum chaperone gene. bZIP is useful for controlling the expression of endoplasmic reticulum chaperone either positively or negatively in cells and therefore is useful for treatment or prophylaxis of ancers, arteriosclerosis, cystic fibrosis, ischaemic diseases, wounds and ulcers. bZIP also maintains the correct conformation of the endoplasmic reticulum chaperone and thereby increases the expression of a foreign protein. This sequence taken from the glucose regulating protein (GRP) promoter GRP94 contains an BRSE like sequence. (Updated on 15-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; protein kinase; PTK; STK; cancer; cardiovascular disease; SNP; metabolic disorder; immune related disease; neurological disorder; neurodegenerative disorder; inflammatory disorder; infectious disease; reproductive disorder; gene therapy; single nucleotide polymorphism; ds
                                                                                                                                                                                                                                                                                                                                          New endoplasmic reticulum stress transcription factor (known as bZIP) fr
controlling expression of endoplasmic reticulum chaperone, useful for
treating cancers, arteriosclerosis, cystic fibrosis, ischemic diseases,
wounds and ulcers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 4 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP containing protein kinase DNA sequence #54.
ulcer; gene therapy; recombinant gene; ci
glucose regulated protein; promoter; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Fig 3; 157pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       150 GAGGCCGGCTTCGACTGG 167
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                                                                                                                                                       99WO-JP006305
                                                                                                                                                                                    98JP-00324227.
99JP-00163112.
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                                                                                                                                                                                                                                          (HSPR-) HSP RES INST INC.
                                                                                                                                                                                                                                                                             Yoshida H,
                                                                                                                                                                                                                                                                                                              WPI; 2000-387736/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                    WO200029429-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                   Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-SEP-2001
                                                                                                                                                                                          13-NOV-1998;
                                                                                                                                                                                                           09-JUN-1999;
                                                                                                                                                         L2-NOV-1999;
                                                                                                                      5-MAY-2000.
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Matches
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AASO6832-AASO6897 represent part of a polymucleotide sequence encoding for novel human protein kinases where a single nucleotide polymorphism (SNP) has been identified. The SNP occurs at the last position of the present sequence. The sequences are described relating to the invention of novel human protein kinases #1-57 (AAU03501-AAU03557). The novel correction kinase (PTK and STK) families. The polymorleotides serine/threonine kinase (PTK and STK) families. The polymorleotides canceding protein kinase and the polymorphies. The polymorleotides prevention, diagnosis and treatment of diseases associated with prevention, diagnosis and treatment of diseases associated with cancers (especially cancers of harmatopic-tic origin), cardiovascular cancers (especially cancers of harmatopic-tic origin), cardiovascular disease (e.g. atherosclerosis), metabolic disorders (e.g. diabetes), immune related diseases (e.g. rheumatoid arthritis), neurological fiscances (e.g. disease), inflammatory disorders (e.g. asthma), infectious disease (e.g. HIV) and reproductive disorders (e.g. asthma), infectious therefore the pranty and as NDA protein kinases may be used for the pharmary and as NDA protein disperse may be used for the pharmary and as NDA protein disperse may and parmoral associated when protein kinase may are parmoral parameters.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy and as DNA probes in diagnostic assays. The protein Kinase polypeptides may be used as antigens in the production of antibodies against the protein kinases and in assays to identify modulators of
                                                                                                                                                                                                                Nucleic acids encoding human kinase polypeptides, useful for preventing diagnosing and/or treating e.g. cancer, immune, cardiovascular and neuronal-associated diseases, and microbial infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Canine; H gene; antiviral; gene therapy; distemper; PCR primer; ss.
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0
                                                                                                                      Sudarsanam S, Martinez R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.1%; Score 13.2; DB 1; Length 19;
83.3%; Pred. No. 3.4e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 2 A; 5 C; 6 G; 5 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Canine distemper virus H gene PCR primer RH-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein kinase expression and activity
                                                                                                                                                                                                                                                                                                 Example 8B; Page 333; 433pp; English
                                                                                                                          Manning G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35 GGACGAAGATGGCCACCA 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 GGCCAAAGATGGCCTCCA 1
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Local Similarity 83.3%;
hes 15; Conservative (
            22-NOV-2000; 2000WO-US032085.
                                                   99US-0167482P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF86572 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUL-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Canine distemper virus.
                                                                                                                            Whyte D,
Clary D;
                                                                                                                                                                                   WPI; 2001-343950/36.
                                                                                     (SUGE-) SUGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2000350587-A.
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                                                   24-NOV-1999;
                                                                                                                              Plowman GD,
                                                                                                                                                Flanagan P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF86572;
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Matches
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for

Yura T;

Mori K, Yanagi H,

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Gaps

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3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02; tive 0; Mismatches 3; Indels

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Mismatches

.; 0

Conservative

15;

Matches

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The invention relates to a method for determining whether an individual is likely to be susceptible to malignant melanoma, and determining the genetic basis for the melanoma in an individual. The method involves screening the genome of the individual for the presence or absence of one or more polymorphic variants of the XRCC3 gene. Sequences AAH47412-420 represent PCR primers used in a genotyping assay of a candidate DNA repair gene XPD
                                                                                                                                     The present invention relates to the H gene derived from canine distemper virus (see AAF86567). The H gene sequence can be used in the prevention, treatment and detection of mammalian distemper, particularly canine distemper virus (CDV). The present sequence is a PCR primer, which was used in the present invention
                                                        H gene, used for treating, preventing and detecting mammalian distemper, particularly canine distemper viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Determining the susceptibility of an individual to malignant melanoma, involves screening the genome of the individual for the presence or absence of one or more polymorphic variants of the XRCC3 gene.
                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  melanoma; genotyping; DNA repair gene; XPD; PCR primer;
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0
                                                                                                                                                                                                                                                               3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 6 A; 4 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                 Sequence 19 BP; 4 A; 6 C; 6 G; 3 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Welsh K;
                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wojnarowska F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       XPD gene exon 23 amplifying primer.
                                                                                                          Example 2; Page 6; 18pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 14; 35pp; English.
                                                                                                                                                                                                                                                                                                                               293 GGTGAAGGACCTGAGCCC 310
                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-FEB-2001; 2001WO-GB000753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-FEB-2000; 2000GB-0004193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS INNOVATION LTD
                                                                                                                                                                                                                                                                                                                                                               GCTGGAGTACCTGAGCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH47419 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 83.33
Matches 15; Conservative
(KYOR-) KYORITSU SHOJI KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Haldar N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-557711/62.
                            WPI; 2001-268280/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymorphism; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200162964-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     XPF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Winsey S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH47419;
                                                                                                                                                                                                                                                                                                                                                                13
                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 295
                                                                                                                                                                                                                                                                                                                                                                                                                         AAH47419
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in method comprises: (b) clones of the genomic libraries contained in method comprises: (b) a primer designed based on the chromosome marker comultiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is defected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell containing the clones in the mixed respecified to a call tured and the resultant cultures are amplified by using the above primer; (g) signals resultant cultures are amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The reconstituted as the positions on the chromosome and arrayed. The represent PCR primers for human chromosome 1956-35 DNA, and ABL45223 to ABL45231 to ABL45231 to C specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                              Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Pred. No. 3.4e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                            Human chromosome 1p36-35 PCR primer SEQ ID NO:1028.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 5 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 25; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTGCTGGCGGCGGACGAC 338
69
                              AATCAGAGGAGACGCTGC 19
                                                                                                                       BB
                                                                                                                                                                                                                                                                                                                                                                                                                              12-MAR-2001; 2001JP-00068285.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-MAR-2000; 2000JP-00066716.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 83.3%;
52 ACTCAGAGGAGTCTCTGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIKA ) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                       ABL43984 standard; DNA; 19
                                                                                                                                                                                            11-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 83.3
les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arraying genome clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-144136/19
                                                                                                                                                                                                                                                                                       PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                           JP2001321190-A
                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                N
                                                                                                                                                           ABL43984;
                                                                                                      ABL43984
                                                                                       RESULT
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GTGCTGGTGGCTGACAAC

13

셤

RESULT 297

Length 19;

3.1%; Score 13.2; DB 1; 83.3%; Pred. No. 3.4e+02;

Query Match Best Local Similarity antiniflammatory steroid; ubiquinone; antiniflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

antisense; lung dysfunction; nasal airway dysfunction;

Human IL4-R oligonucleotide sequence.

(first entry)

17-OCT-2003

ABZ97333;

ABZ97333 standard; DNA; 19

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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                Human, antisense, lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiantsthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 12494; 872pp; English.
                                                                                                                                              adenosine receptor; bronchodilation; bronch
lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                Li Y, Sandrasagra A,
Tang L, Shahabuddin
           ABZ97252 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                      24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                          EPIG-) EPIGENESIS PHARM INC
                                                                             Human nucleic acid sequence.
                                                      17-0CT-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-229219/22.
                                                                                                                                                                                                    WO200285308-A2.
                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                         31-OCT-2002.
                                                                                                                                                                                                                                                                                                                Nyce JW, 1
                                                                                                                                                                                                                                                                                                                                                                                                       ubiquinone
                                  ABZ97252;
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intitation coodon, coding regalon, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entithflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antientlammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of, or reducing sensitivity to adenosine, reducing levels of, or reducing sensitivity to adenosine, reducing levels of ubiquinone or lung surfactant in a subject's tissue, or treating pronchoorseriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed appearance in the printed appearance in the printed appearance in the printed appearance of the sequence data for this patent is not represented in the printed appearance in the printed appearance for this form when the printed and the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 2 A; 10 C; 3 G; 4 T; 0 U; 0 Other;
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an continifiammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a comparing a respiratory, lung or mallgnant crespicatory effect of an continifiammatory steroid in a subject, for reducing or depleting levels of or entiantflammatory steroid in a subject, for reducing or depleting levels of continifiammatory steroid in a subject, for reducing levels of adenosine continity producing bronchodilation, increasing levels of adenosine contriblemmation, lung allergies, or treating bronchoconstriction, clung inflammation, but was obtained in electronic format directly from WIPO at the subject is a respiratory disease or condition.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

Disclosure; SEQ ID NO 12575; 872pp; English.

ubiquinone.

Katz E, Pabalan J, Aguilar D;

Li Y, Sandrasagra A, Ka , Tang L, Shahabuddin S;

WPI; 2003-229219/22.

Miller S,

Nyce JW,

Aguilar D;

Katz E, Pabalan J, S;

23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC

WO200285308-A2 Homo sapiens.

31-OCT-2002.

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at ftp.wipo.int/pub/published_pct_sequences
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Gaps
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Pred. No. 3.4e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 2 A; 10 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13
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Best Local Similarity 83.3%;
Matches 15; Conservative C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      337 ACCAGGGCCGGCTGCTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACCACGCCCGGCTTCTCT
ABZ97333
ABZ973333
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Gaps

3.1%; Score 13.2; DB 1; Length 19; 83.3%; Pred. No. 3.4e+02; ve 0; Mismatches 3; Indels

83.3%;

15; Conservative

Similarity

Local Best Loca Matches 337 ACCAGGCCGGCTGCTCT 354

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ACCACGCCCGGCTTCTCT

RESULT 298

at ftp.wipo.int/pub/published\_pct\_sequences

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RESULT 299

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WPI; 2003-268345/26.
                                                                                     misc_difference 19
                                           Synthetic.
Hepatitis C virus.
                                                                                                                              WO2003016572-A1.
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07-JJL-1992
                                                                                                                                                   27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ22593;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           stem_loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 301
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ID AAQ22
XX
   THXSXEXBXBXBXE
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                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated double stranded RNA oligomucleotide about 19 to about 25 ribonucleotides in length or its equivalent. One strand of the oligomucleotide comprises the same nucleotide sequence as a region of a hepatitis C virus (HCV) target polymucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligomucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the anti-HCV agent LZ129 mutant RNA of the invention which contains a C3G mutation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                /*tag= a
/note= "Wild-type cytosine substituted for guanine"
                                                                                                                                                                                                                                                                                                                                                                                    New double stranded RNA oligonucleotide, useful for preparing a composition for treating or preventing hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ô
                                                                                            HCV infection; replication; pathogenesis; virucide; vaccine; gene therapy; ds; anti-HCV; agent LZ129; mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.2; DB 1; Length 19; 72.2%; Pred. No. 3.4e+02; tive 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3 A; 6 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                           Zhao G, Lu J, Glass JI, Martinez A, Yang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-HCV agent LZ129 mutant RNA - G4C.
                                                                         Anti-HCV agent LZ129 mutant RNA - C3G.
                                                                                                                                                 Location/Qualifiers
misc_difference 19
                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 155; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    256 CGGCCACGGTGCACCTGG 273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CGGGCACGAUGCAUCUGG 18
          ADD00872 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                           17-AUG-2001; 2001US-0313076P.
20-DEC-2001; 2001US-0344116P.
01-FEB-2002; 2002US-0353750P.
                                                                                                                                                                                                                                                         16-AUG-2002; 2002WO-US021843.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD00871 standard; RNA; 19
                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3.1%
Query Match
Best Local Similarity 72.2%
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-268345/26.
                                                                                                                            Synthetic.
Hepatitis C virus.
                                                                                                                                                                                                               WO2003016572-A1
                                                     01-JAN-2004
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                                                                                                                                                                                                                                    27-FEB-2003
                                ADD00872;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD00871;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 300
ADD00872
ID ADD0
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The invention relates to a novel isolated double stranded RNA oligonucleotide about 19 to about 25 ribonucleotides in length or its equivalent. One strand of the oligonucleotide comprises the same nucleotide sequence as a region of a hepatitis C virus (HCV) target RNA polynucleotide sequence required for hepatitis C virus infection, replication or pathogenesis in vitro or in vivo in a host cell. The oligonucleotide of the invention demonstrates virucide activity and may be useful for preparing a composition or vaccine for treating or preventing hepatitis C virus, as well as during gene therapy procedures. The current sequence is that of the anti-HCV agent LZ129 mutant RNA of the invention which contains a G4C mutation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                          /*tag= a
/note= "Wild-type guanine substituted for cytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New double stranded RNA oligonucleotide, useful for preparing a composition for treating or preventing hepatitis C virus.
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HCV infection; replication; pathogenesis; virucide; vaccine; gene therapy; ds; anti-HCV; agent LZ129; mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.1%; Score 13.2; DB 1; Length 19; Best Local Similarity 72.2%; Pred. No. 3.4e+02; Matches 13; Conservative 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3 A; 8 C; 5 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhao G, Lu J, Glass JI, Martinez A, Yang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 155; 173pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CGCCCACGAUGCAUCUGG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-AUG-2001; 2001US-0313076P.
20-DEC-2001; 2001US-0344116P.
01-FEB-2002; 2002US-0353750P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-AUG-2002; 2002WO-US021843.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ELIL ) LILLY & CO ELI
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/\*tag= a /note= "forms loop with substrate" /note= "forms loop with substrate" /\*tag= b /tag= b

misc\_RNA

Guerrierta CL

Forster AC,

UYYA ) UNIV YALE. .7-AUG-1990;

IPI; 1992-096909/12

91WO-US005808 90US-00568834

W09203566-A

Disclosure; Fig 2d; 34pp; English

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To develop antisense oligos, the test system employed mouse NIH 3T3 cells stably transfected with an internally deleted construct of the human gene for the pro alpha 1(1) chains of type I procollagen COLLAI). A series of modified oligos were synthesised using a region at the 3' end of exch 1 and the first two nucleotides of intron 1 of the exogenous (human) gene as a target. This sequence is given in AAQ66555 which corresp. to bps 198 -225 if the adenine at the start of transcription is counted as poon. The corresp, which corresp to bps 169 199 months and the first the adenine at 10 pps 169 195 months are sequence of the endogenous (mouse) gene is given in AAQ66557, which corresp to bps 169 199. The antisense oligos are jiven in AAQ665597-Q66614. The antisense oligos inhibit the expression of mutant or normal collagen genes. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                             Anti:sense oligo:nucleotide(s) against mutant or native collagen genes for inhibiting collagen expression, e.g for treating osteoarthritis, liver cirrhosis, excessive scarring etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human type I procollagen (COLIAI) pro alpha 1 chain antisense
oligonucleotide AS9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 7 C; 5 G; 8 T; 0 U; 0 Other;
Procollagen; antisense oligo; inhibition; ss.
                                                                                                                                                                                               Nugent P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Procollagen; antisense oligo; inhibition;
                                                                                                                                                                                             Baserga R,
                                                                                                                                                                                                                                                                                                            Claim 5; Page 24; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26 CGAGGGCTGGGACGAAGA 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ66602 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                    (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 CGAGGGCCAAGACGAAGA
                                                                                                                                         92US-00973332.
                                                                                                              93WO-US010756
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(first entry)
                                                                                                                                                                                               Prockop D, Colige A,
                                                                                                                                                                                                                         WPI; 1994-183496/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
10-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-MAY-1994
                                                                                                                                         09-NOV-1992;
                                                        WO9411494-A1
                                                                                   26-MAY-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ66602;
                             Synthetic.
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AAQ66602/c
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0
                                                                                                                                                                                                                                                                                 Compsn. for targeting RNA sequence for cleavage by RN ase P - comprises external guide sequence including 3-NCCA and complementary nucleotide sequences, for treating viral diseases.
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Gaps . 0

Length 20;

Score 13.2; DB 1; Length 20 Pred. No. 3.8e+02; 0; Mismatches 3; Indels

Query Match
3.1%;
Best Local Similarity 83.3%;
Matches 15; Conservative

243 TGCTTCCCGGGCTCGGCC 260

N

19 TGGTGCCCGGACTCGGCC

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Seguence 20 BP; 3 A; 8 C; 8 G; 0 T; 1 U; 0 Other;

Human type I procollagen (COLIAI) pro alpha 1 chain antisense oligonucleotide AS8.

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Nugent P;

Baserga R,

Colige A,

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To develop antisense oligos, the test system employed mouse NIH 373 cells stably transfected with an internally deleted construct of the human gene for the pro alpha 1(1) chains of type I procollagen CCLIAI). A series of modified cligos were synthesised using a region at the 3' end of exon I and the first two nucleotides of intron 1 of the exogenous (human) gene as transpet. This sequence is given in AAG66555 which corresp. to bys 198 198 -225 if the adenthe at the start of transcription is counted as poon. The corresp, sequence of the endogenous (mouse) gene is given in AAG66559, which corresp to bys 169-195. The antisense oligos are given in AAG66559. Goodeld. The antisense oligos inhibit the expression of mutant or normal collagen genes. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A novel DNA construct (preferably a retroviral vector) has been produced for the treatment of human mammary cell disorders or diseases, including human mammary carcinoma. The DNA construct comprises at least one therapeutic gene under the transcriptional control of the whey acidic
                                                                Anti:sense oligo:nucleotide(s) against mutant or native collagen genes for inhibiting collagen expression, e.g for treating osteoarthritis, liver cirrhosis, excessive scarring etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene expression; human mammary carcinoma cell; whey acidic protein; mouse mammary tumour virus; WAP; WMTV; polymerase chain reaction; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression in human mammary carcinoma cells - using whey acidic in or mouse mammary tumour virus regulatory sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine leukaemia virus retroviral vector BAG PCR primer B.
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 8 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BAVA-) BAVARIAN NORDIC RES INST AS.
(GSFU-) GSP FORSCHUNGSZENTRUM UMWELT & GESUNDHEI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Fig 1; 46pp; English.
                                                                                                                                             Claim 5; Page 24; 55pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26 CGAGGGCTGGGACGAAGA 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT62029 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18 cgadddccaagacgaaga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein or mouse mammary
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 83.3
nes 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-192915/17.
                                 WPI; 1994-183496/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9709440-A1
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14-NOV-1997
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Prockop D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT62029;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 304
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(first entry)

(revised)

Saller RM, Salmons B;

95DK-00000976. 96WO-EP003922.

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A retroviral vector carrying a DNA sequence encoding SDI-1 (senescent cell derived inhibitor 1), a functional analogue, fragment or antisense SDI-1 DNA sequence has been developed. The present sequence represents FOT primer B used in the amplification of mouse leukaemia virus (MLV) retroviral vector beta galactosidase gene (BAG) LTR, for use in the cetroviral vector can be used in the treatment of a polylinker. The retroviral vector can be used in the treatment of disorders or diseases responsive to the anti-proliferative activity of SDI-1, e.g. for the treatment of cencer or restenosis, sepecially for the treatment of breast cancer. The retroviral vector acts to introduce the relevant DNA sequences, sense or antisense, into human cells in vitro or in vivo. The retroviral vector antistered by injection or by implantation of a packing cell ine in to the body nearby or at the site of the tumour. (Updated on 25-MAR-2003 to correct PI field.)
The present sequence represents PCR primer B which is involved in the deletion of the U3 region from the murine leukaemia virus (MLV) retroviral vector, known as BAG, and the insertion of a polylinker, which is used in an example for the production of a DNA construct as described above. The WAP and MMTV regulatory sequences are able to direct the mammary calls, including mammary carcinoma cells. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Retroviral vector carrying senescent cell derived inhibitor 1 DNA - used in the treatment of diseases responsive to anti:proliferative activity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MLV; retroviral; vector; senescent cell derived inhibitor 1; SDI-1; antiproliferative; breast cancer; restenosis; human; implantation; tumour; polymerase chain reaction; beta galactosidase gene; ss.
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3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.88+02;
Matches 15; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BAVA-) BAVARIAN NORDIC RES INST.
(GSFU-) GSF FORSCHUNGSZENTRUM ITMY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT85369 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-OCT-1995;
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0; Gaps

3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels

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Gaps

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Lead in the chimeric non-human animal of the invention. The chimeric non-human animal of the invention. The chimeric non-human animal of the invention, preferably a mouse, contains a foreign chromosome (s) or chromosome fragment. The animal is produced by obtaining a hybrid cell by fusion of a cell containing the foreign chromosome with a repared, and fused with cells having differentiative pluripotency and containing the promosome. These cells are then introduced into an embryo, which is then chromosome. These cells are then introduced into an embryo, which is then chromosome segment to term. The foreign chromosome segment is at least in the long and preferably contains a region for an antibody. The chromosome segment could also contain genes associated with human chromas is useful for efficient production of proteins, especially thuman animal is useful for efficient production of proteins, especially of human animal is useful for efficient production of proteins, especially of thuman animal is useful material calls of the chimeric animal which myeloma cells to produce hybridomas capable of expressing the foreign gene (e.g. to produce the antibody)
                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer; amplify; human gene; chimeric non-human animal; antibody; transgenic mouses; chromosome fragment; hybridoma production; microcell; Huntington's disease gene; pluripotent cell; interleukin-2 gene; myeloma cell; immunoglobulin gamma-1; constant region; IGG1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric animal containing foreign chromosome - for expression of a foreign gene, e.g. an antibody.
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                                                                                                                                                                                                                                                                                                                                                                          Primer #2 for immunoglobulin gamma-1 constant region (IGG1).
                                                 Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanaoka K, Oshimura M, Ishida I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
               Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 9; Page 33; 142pp; Japanese.
                                                                                                                               47 CCACCACTCAGAGGAGTC 64
                                                                                                                                                                 AAT92797 standard; DNA; 20 BP.
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96JP-00027940.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96WO-JP002427.
                                                   Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-178822/16.
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The present invention describes a method of obtaining pluripotent cells containing foreign chromosomes or their fragments (preferably at least containing foreign chromosomes or their preparing cancerous cells containing the foreign chromosomes or fragments, then fusing these with pluripotent cells such as embryonic stem cells, embryonic reproductive cells embryonic cancer cells or their mutants. Also cascribed are: (1) a method of obtaining hybridoma cells (such as mouse A9 cells) with a high ability to produce hybridoma cells (such as mouse A9 cells) with a cell containing the foreign chromosomes or fragments (such as normal human diploid cells); (2) a method of utilising pluripotent cells or produce chimeric and transgenic non-human animals (especially mammals such as mice) which can express the foreign chromosomes or fragments introduced; and (3) chimeric animals, their offspring and tissues and cells derived from the offspring produced by a method as in (2). The introduced; and (3) chimeric animals, their offspring and tissues and cells are of human type and therefore not antigent in humans. They can also be used in the production of chimeric and serve as models for the study of human diseases. Adv52755 to A4V52828 are perve as models for the study of human diseases. Adv52755 to A4V52828 are
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                                                                                                                                                                                                                                                                                                                                                              construction; human; antibiotic gene; cancer cell; embryonic; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pluripotent cells containing foreign chromosomes or fragments - and n
human chimeric animals constructed using them and expressing foreign
genes such as human antibiotic genes.
                    Gaps
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                                                                                                                                                                                                                                                                                                                                                Pluripotent cell; intrinsic gene; chimeric non-human animal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hanaoka K, Oshimura M, Ishida I;
83.3%; Pred. No. 3.8e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                             Immunoglobulin gamma-1 constant PCR primer IGG1 #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 9; Page 46; 217pp; Japanese.
                                                           364 TCCTCACTTTCCTGGACC 381
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-JP000860.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97JP-00062309.
                                                                                                20 recreaceatecaeee
                                                                                                                                                                                              AAV52794 standard; DNA; 20
      Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fomizuka K, Yoshida H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-480821/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-1997;
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                                                                                                                                                                                                                                                                          27-NOV-1998
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                                                                                                                                                                                                                                   AAV52794;
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DB 1; Length 20;

3.1%; Score 13.2;

Query Match

AAZ25788 standard; DNA; 20 BP.

AAZ25788/c

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This sequence represents Basta-resistance gene (bar) PCR primer Bar 2, used with primer Bar 1 (AAZ12671) to determine whether the bar gene had been stably introduced into the genome of Basta-resistant plantlets. The plantlets had been previously been transformed with a superoxide plantlets had been previously been transformed with a superoxide of promoter (SOD) expression vector comprising a cucumber fruit dominant promoter (SOD), a cassava mSODI gene and the bar gene. SODs are ubiquicous enzymes which convert superoxide anion radicals to hydrogen peroxide anion radicals, and other reactive oxygen species such as hydrogen peroxide and hydroxyl radical, are generated by cranisms in response to environmental and biological stresses. SOD is thought to be effective in the treatment of arthritis, rheumatism, is chaemic heart disease and radiation damage, and has been used in commetics for the prevention of skin ageing? The transgenic plants (especially cucumber fruit) produced by the method of the invention are used as materials for cosmetics in medicines. The plants can be used as materials and a series of medicines. The plants can be used as
                                                                                                                                                                                                                                                                 SOD; superoxide dismutase; antioxidant; superoxide; anion; radical; environmental stress; biological stress; treatment; arthritis; rhemunatism; ischaemic heart disease; radiation damage; skin; ageing; cosmetic; plant bioreactor; transgenic plant; expression; herbicide; Basta; resistance; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Producing transgenic plants which produce high levels of superoxide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                Basta-resistance (bar) gene PCR primer Bar 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (KOAD ) KOREA ADV INST SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kim J, Lee H, Kwon SY, Kwak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98KR-00013205.
98KR-00033947.
99KR-00011848.
                                                                                                                  BP.
20 TCCTCACCGTCCTGCACC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99EP-00302909.
                                                                                                                    AAZ32672 standard; DNA; 20
                                                                                                                                                                                              09-FEB-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Basta-resistant plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-582804/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-AUG-1998;
06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                           AAZ32672;
                                                                               RESULT 308
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                                   Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
Sequence 20 BP; 1 A; 4 C; 6 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                   Mouse ss3 gene reverse PCR primer.
                                                                                                                      181 CCAAGGCACATATCCACT 198
                                                                                                                                                            20 czasedczeczace 3
                                                                                                                                                                                                                                                               AAV64430 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                               01-MAR-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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Gaps

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3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; cive 0; Mismatches 3; Indels

214 AGAACTCGGTGGCGGCCA 231

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15; Conservative

Local Similarity

20

AGATCTCGGTGACGGGCA

RESULT 309

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The present sequence represents a PCR primer for the human p51 gene, which is related to p53 and has cell proliferation regulation and tumour suppression activity. The p51 gene can be used in the investigation, diagnosis and treatment of diseases such as cancer, with which the p53 family cell proliferation regulation is associated. The p51 protein may be used for screening potential agonists and antagonists of its regulatory function, for use as drugs,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          se3 gene; mouse; liver development; signal transduction; liver disease;
tissue repair; cancer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                    New p53 related human gene p51, useful for diagnosis, investigation and treatment of cancers and screening for potential cell proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                               Human, p51, p53 related gene, cell proliferation, regulation, cancer;
tumour suppression, diagnosis, PCR primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 116; 163pp; Japanese.
                                                                                                                                                                                                                                                                                       Obinata M;
                                                                                                                                                                                                                              98JP-00100467.
                                                                                                                                                                                                     99WO-JP001512.
                                                                                                                                                                                                                                                (SAKA ) OTSUKA PHARM CO LTD.
(IKAW/) IKAWA Y.
                                                          Human p51 PCR primer p51-R6.
                                   (first entry)
                                                                                                                                                                                                                                                                                                               WPI; 1999-591318/50.
                                                                                                                                                                                                                                                                                       Ikawa Y, Ikawa S,
                                                                                                                                                                                                       24-MAR-1999;
                                                                                                                                                                                                                              27-MAR-1998;
                                                                                                                                Homo sapiens
                                                                                                                                                       WO9950412-A1
                                   07-JAN-2000
                                                                                                                                                                                07-OCT-1999.
                                                                                                                      Synthetic.
              AAZ25788;
                                                                                                                                                                                                                                                                                                                                                                 agents.
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05-NOV-1998

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PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antiense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, oppidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and veneral lymphogranulomatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccine, eye disease, conventional trachoma, nonendemic trachoma, paratrachoma, inclusion conjunctivitis, genital disease, perihepatitis, nongonococcal uretritis, epidymitis, cervicitis, salpingitis, PCR primer, bartholinitis, pneumopathy, venereal lymphogranulomatosis, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFS) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFS encode polypeptides (see AAY36794-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genome sequence of Chlamydia trachomatis.
                Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 1655; 1755pp; English.
                                                               Disclosure; Page 1600; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               351 CTCTACAGCGACTTCCTC 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 crccacaccaarrerre 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97FR-00015041.
97FR-00016034.
98US-0107077P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ04026 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-371125/31.
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04-NOV-1998;
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AAZ04026/C
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                                                                                                                                                                                                                                                                                                                                                                                        This oligonuclectide is used as a reverse primer, together with a forward primer (see AAV64429), in quantitative PCR of the mouse ss3 gene. 10 demonstrate pairs (see AAV64427-46) were used in quantitative PCR to demonstrate elf (see AAV644113), sa3, and 145 (see AAV64414) expression in bloc and liver explant cultures, compared to HNF3-beta, C/EBP, alphafetorotein and glyceraldehyde 3-phosphate dehydrogenase. The invention provides early developing liver proteins and the genes coding for them (see AAV64410-24). These can be used in the treatment and diagnosis of liver diseases and other disorders, including those relating to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine, eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                              New isolated early liver development genes - used to develop products for treating, e.g. liver disease, hepatocellular carcinoma, degenerative neurological disorders, anaemia, ataxia or haemochromatosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                          Example 2; Fig 20; 92pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 347 GCTGCTCTACAGCGACTT 364
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97FR-00016034.
98US-0107077P.
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                                               98WO-US008656.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.1%;
                                                                                         97US-00841349
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Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
Chlamydia trachomatis.
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                                                                                                                                   MISH/) MISHRA
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04-NOV-1998;
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                                                                                         30-APR-1997;
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RESULT 31

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WPI; 1999-371125/31

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Gaps

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BP.

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Human D2 dopamine receptor PCR forward primer.
AAX00253 standard; DNA; 20
                                                               (first entry)
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Best Local Similarity 83.3
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     Kaddis FG;
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-095293/08.
                                                                                                                                                                                          Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                        WO9857663-A1.
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                                                               26-MAR-1999
                                                                                                                                                                                                                                                                        23-DEC-1998.
                                                                                                                                                                                                                                                                                                                                                                                                     Freed CR,
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                                AAX00253;
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AAZ98749/c
 88XCCCCCCCCCCX8XEXEXEXEXEXEXEXEXEXXEXXEXXEXXEXXEX
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conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococoal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents a primer/probe sequence of the invention. The primer and probe sequences are derived from the sequence of the human serine protease gene LKB1, and are used to detect variations in LKB1 leading to Peutz-Jeghers (PD) syndrome. The primers and probes can be used for the diagnosis, investigation and treatment of diseases in which variations in the LKB1 gene are implicated, such as PJ syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                   LKB1 gene; human; serine protease; Peutz-Jeghers syndrome; PJ syndrome; variation detection; therapy; diagnosis; primer; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primers and probes for use in diagnosis of Peutz-Jeghers syndrome
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                                                                                                                                        Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; les 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 6 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                             Sequence 20 BP; 1 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 89; 107pp; Japanese
                                                                                                                                                                                                           225 GCGGCCAAATCGGGAGGC 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        89
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                                                                                                                                                                                                                                          GCTGCCAAAGCGGGAGCC 3
                                                                                                                                                                                                                                                                                                                        AAX79655 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-JP005357.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97JP-00344256
98JP-00280357
                                                                                                                                                                                                                                                                                                                                                                                                                      Human LKB1 gene primer/probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        51 CACTCAGAGGAGTCTCTG
                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nezu J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W09928459-A1
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                                                                                                                                                                                                                                                                                                                                                                                       12-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                         AAX79655;
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Best Local
                                                                                                                                             Query Match
                                                                                  diseases.
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                                                                                                                                                                              Matches
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RESULT AAX7965

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A method has been developed of treating defective tissue comprising: (i) providing a number of hWT neurons and a neurologically defective mammal having a target trissue comprising defective cells, and (ii) transplanting the hWT neurons into the defective mammal so that the neurological defective for the mammal is ameliorated. Also described is a non-human mammal having transplanted hWT neurons. The method is especially used to treat thutington's disease or other neurological disorders. The method allows the transplantent of terminally differentiated neurons from call lines. The present sequence represents a PCR primer used in an example from the present invention for in vitro characterisation of hWT neurons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             - by
Human; glutamic acid decarboxylase; choline acetyltransferase; GAD65; GAD67; ChAT; dopamine receptor; G3PDH; PCR primer; Huntington's disease; neural transplantation; neurological disease; hVT neuron; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treatment of neurological disorders, especially Huntington's disease transplantation of differentiated neurons into corpus striatum of affected mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; mitochondria; PCR primer; large insert episome; lipofection; epstein barr virus nuclear antigen-1; BBNA-1; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.1%; Score 13.2; DB 1; Length 20;
83.3%; Pred. No. 3.8e+02;
tive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 36; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  289 AGCTGGTGAAGGACCTGA 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYTE-) UNIV TECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                               98WO-US012685.
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RESULT 314 AAX00253/c

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99WO-US019468. 98US-0097961P. 98US-0102691P.

6-AUG-1999;

09-MAR-2000.

01-OCT-1998;

26-AUG-1998;

WO200012693-A1

UYNC-) UNIV NORTH CAROLINA

WPI; 2000-256638/22.

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                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is that of nested primer NP2, which was used in the amplification of gene fragments obtained from a suppression subtractive hybridization reaction using LAPC xenograft cDNA and designed to identify novel prostate and prostate cancer-specific genes. A 437 bp clone was obtained. Pull-length cDNA (see AAZ94275) was subsequently cloned from a transcription factor that is normally expressed only in testis tissue, but is up-regulated in prostate and other types of cancer. The invention provides diagnostic and therapoutic methods useful in the management of various cancers which express PHELIX, including prostate cancer, bladder cancer, ovarian cancer and testicular cancer
                                                                                                                                                                                                                                                         Testis specific Helix Loop Helix proteins expressed in cancers and useful for the prevention, diagnosis and treatment of prostate, bladder and ovarian tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumour necrosis factor receptor-associated factor; TRAP; human;
antisense oligonucleotide; phosphorothioate; antiproliferative;
anti-inflammatory; B-selectin; jun kinase; ss.
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3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TRAF2 antisense oligonucleotide ISIS# 16847.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Xu XS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP,
                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 31; 62pp; English
                                                                                                                                                                             Raitano AB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 recreseceses 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAASSSS6 standard; DNA; 20 BP
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98US-0098610P.
98US-0106524P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                UROGENESYS INC
                                                                                                                                                                             Afar DE, Hubert RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-303732/26.
                                                                                     AFAR D E.
HUBERT R S.
RAITANO A B.
                                                                                                                                                                                                                      WPI; 2000-237872/20
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31-AUG-1998;
31-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA55556;
                                                                                     (AFAR/)
(HUBE/)
(RAIT/)
                                                                UROG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                               this sequence represents a PCR primer used to amplify a fragment of the human mitochondrial DNA. The PCR product is used to create a probe which is used in the Southern blot analysis of cells transfected with the recombinant plasmid of the invention. The plasmid is useful for the production of large-insert episomes in mammalian cell. The plasmid comprises a lymphotrophic herpes virus segment and a heterologous insert cepitation and a heterologous origin of plasmid replication and a heterologous origin of bacterial replication which is maintained as an episome in both bacterial and mammalian cells, cepicially B-lymphoblastoid cells (BLC), epithelial cells (BC) or a ceombinant plasmid is useful for transforming mammalian cells, specially B-lymphoblastoid cells (BLC), epithelial cells (BC) or a fusion of these, by transfecting a mammalian cell with the plasmid by lipofection. The recombinant plasmid is also useful for the production of large-insert episomes in mammalian cells. The invention also relates to a repetien barr virus nuclear antigen-1 (BBNA-1) gene having a partial IR3 domain deletion which is from 300-700 nucleotides in length. The BBNA-1 energy gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                 New recombinant plasmid useful for producing large-insert episomes in mammalian cells comprises a lymphotrophic herpes virus segment linked to a heterologous insert segment.
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Gaps ;

3.1%; Score 13.2; DB 1; Length 20; llarity 83.3%; Pred. No. 3.88+02; Conservative 0; Mismatches 3; Indels

Best Local Similarity Matches 15; Conserv

Query Match

CTGCTCGGTGAAAGCAGA 214

197

8

0

CTGCTAGGTGTAAGGAGA

19

ВР

AAZ94278 standard; DNA; 20

RESULT 316

Human PHELIX nested primer NP2

03-JUL-2000 (first entry)

AAZ94278;

99WO-US020137.

31-AUG-1999;

09-MAR-2000

WO200012709-A2.

Homo sapiens

Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;

Example 2; Page 33; 67pp; English.

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The present invention relates to antisense oligonucleotides (see AAA55496 + A55757) which are targeted to nucleic acids encoding a human tumour necrosis factor receptor-associated factor (TRAP). The antisense sequences comprise at least one modified internucleotide linkage, which is a phosphorothhoate linkage. The oligonucleotides also include at least one modified sugar moiety come modified sugar moiety.

Sequences AAA55490-A55495 represent nucleotide sequences encoding human TRAF1-6. Included in the invention is a method for treating a human having a disease associated with the expression of TRAF comprising administering an antisense oligonucleotide. The reduction of jun kinase activation in cells comprises contacting the cells with an antisense oligonucleotide encoding the cells or tissues with an antisense oligonucleotide targeted to TRAF-6. Method for the reduction of E-selectin expression in cells or tissues comparises contacting the cells or tissues with an antisense oligonucleotide targeted to TRAF-6.
                                                                                                                                                                                                                                                                                                            The antisense oligonucleotides have antiproliferative and anti-
inflammatory activity and are useful for treating disorders associated
with cell proliferation and inflammation. The antisense oligonucleotides
may also be used as a diagnostic probe for studying gene function
diseases associated with TRAF expression such as inflammatory diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PTAN; testis specific; prostate cancer; overexpress; chromosome 1q22; diagnose; cancer; breast; vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PTAN proteins, and sequences encoding them, used for diagnosing and treating cancers, especially breast and prostate cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                   3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 20 BP; 5 A; 9 C; 5 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer (NP2) used in PTAN gene isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                   Example 16; Page 52; 170pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Raitano AB,
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98US-0102910P.
98US-0113239P.
99US-0129518P.
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HUBERT R S.
RAITANO A B.
MITCHELL S C
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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21-DEC-1998;
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(HUBE/)
(RAIT/)
(MITC/)
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Gaps

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                        This sequence represents a PCR primer used in the isolation of cDNA fragments of the PTAN (testis specific protein expressed in prostate cancer) gene. FTAN is expressed in 3 isoforms PTAN-1, 2, and 3. The PTAN gene is located on chromosome 1022. PTAN is overexpressed in prostate cancer, and has a testis specific expression pattern in adult tissues. FTAN shows no homology to any known gene. PTAN can be used in methods for the diagnosis of cancer, especially prostate or breast cancer, where the normal tissue samples are prostate tissue, or breast tissue, bone tissue, lymphatic tissue, serum, blood, or urine. A vector containing the PTAN nucleotide sequence, a vaccine composition targeting PTAN, PTAN, ribozymes specific for PTAN mRNA and antisense sequences, capecially breast and prostate cancers. Cancer development can be inhibited by a vaccine composition targeting PTAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated BPC-1 polypeptides, useful for developing products for the diagnosis, staging, prognosis and treatment of cancers, particularly prostate or bladder cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ogene; oncogenic; cancer; prostate; bladder; antibody; vaccine; detection; prognosis; drug screening; primer; ss
                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer used for generating human brain specific protein BPC-1 cDNA.
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                                                                                                                                                                                                                                                                                                                                  3; Indels
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                                                                                                                                                                                                                                                                                               3.1%; Score 13.2; DB 1; Length 83.3%; Pred, No. 3.88+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                               Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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Example 1; Page 31; 71pp; English.
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                                                                                                                                                                                                                                                                                                                                                                  373 TCCTGGACCGCGACGACG 390
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                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RAITANO A B.
SAFFRAN D C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (LEON/) LEONG K.
(RAIT/) RAITANO A B.
(SAFF/) SAFFRAN D C.
(JAKO/) JAKOBOVITS A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-206006/18.
                                                                                                                                                                                                                                                                                                             Local Similarity
les 15, Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AFAR D E.
HUBERT R S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BPC-1; oncogene;
antisense; vacci
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-AUG-1999;
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(HUBE/)
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AAZ94898/c
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polymucleotides can be used for treating such cancers. The BPC-1 collypeptides can also be used in vaccines for treating or inhibiting the development of a cancer expressing BPC-1. The polypeptides and collypeptides and produced can also be used for detection, prognosis, drug screening comprises a CUB domain which is expressed in prostate and bladder comprises a CUB domain which shows sequence similarity with CUB domains from carcinoma cells and which shows sequence similarity with CUB domains from other known proteins. In normal human tissues BPC-1 is only expressed in prostate cancer cells and bladder cancer cells. A number of synthetic oligonucleotides were used to generate BPC-1 cDNA from total cell RNA of tumour cells lines. These primers were a cDNA synthesis primer (AAZ93041), two adaptor sequences (AAZ93042-Z93045), a PCR primer (AAZ93046) and two nested primers (AAZ93047, AAZ93048). This sequence is one of the nested primers (NP)1 used in the amplification method
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a method for obtaining a transgenic plant with male sterility. The method uses site specific recombination to stably transform the plant cells. The method involves the use of DNA encoding the histocyte lethal protein, linked to an auther specific promoter. The method is used to produce male sterile plants. Sequences AAA59949 to AAAS9961 are used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polynucleotide SEQ ID 13 used in method to culture sterile male plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Male sterile plant; transgenic plant; histocyte lethal protein; anther specific promoter; ss.
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                                                                                                                                                                                                                                                                                                                                                                       Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Molecular method for culturing male sterile plant.
                                                                                                                                                                                                                                                                                                                                  Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 23; 32pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                          373 TCCTGGACCGCGACG 390
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA59961 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hu Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-400684/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Li L'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20~OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel testes-specific gene 22P4F11 which is expressed in human prostate cancer and is useful as a diagnostic marker and/or therapeutic target for prostate cancer.
                                                                                                                                                                                                                            22P4P11; human; testis; prostate cancer; diagnosis; gene therapy; marker; vaccine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expressing 22P4Fil, especially prostate cancer, are provided, as well as vaccines that prevent development of such cancers
                                                                                                                                                                         PCR primer NP2 used in testis-specific 22P4F11 gene amplification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hubert RS, Mitchell SC;
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AAZ94898 standard; DNA; 20 BP
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99US-0146584P.
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Best Local Similarity 83.3
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AFAR/) AFAR D E.
(HUBE/) HUBERT R S.
(MITC/) MITCHELL S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-303452/26.
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                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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28-JUL-1999;
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                                                           AAZ94898;
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ID AAA1
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WO200010383-A1.

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PCR primers AAA14805-07 were used to amplify testis-specific protein Yencoded DNA. The specification describes a new method of diagnosis of prostate cancer. The method comprises determining the level of testis-specific protein Yencoded (TSPY) mRNA or protein, and comparing these TSPY mRNA or protein is indicative of prostate presence of elevated TSPY mRNA or protein is indicative of prostate cancer. Detection of TSPY mRNA expression or protein levels is useful in the diagnosis of prostate cancer. Antisense polymucleotides complementary to the coding sequence of human TSPY are useful for treating prostate cancer by inhibiting TSPY transcription (when contacted with the TSPY mRNA). Ribozymes are also useful for treating prostate cancer by cleaving the TSPY mRNA and therefore inhibiting its translation. The vaccine is useful for inhibiting the development of prostate cancer in a patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Foreign chromosome; microcell fusion; homologous recombination; antibody; targeting vector; transgenic animal; disease model; knockout animal; PCR primer; human; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Diagnosing prostate cancer by determining the level of testis-specific protein Y-encoded (TSPY) mRNA or protein and comparing these TSPY mRNA or protein levels to those of a normal tissue sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer 2 for human immunoglobulin gamma-1 constant region gene IGG1.
                                                                                        Prostate cancer; testis-specific protein Y-encoded mRNA; TSPY mRNA; vaccine; PCR primer; ss.
                                                              PCR primer for testis-specific protein Y-encoded DNA.
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Best Local Similarity 83.39
----hes 15; Conservative
                               (first entry)
                                                                                                                                                                                                                                                                                                                (UROG-) UROGENESYS INC. (AFAR/) AFAR D E.
                                                                                                                                                                                                                                                                                                                                                                    Hubert RS;
                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-303803/26.
                                                                                                                                                                                 WO200020638-A2
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                              08-AUG-2000
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AAA14807;
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AAA09957/c
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The invention relates to a novel method of producing cells containing a modified foreign chromosome or chromosome fragment. The method comprises:

(a) fusing a microcell comprising the foreign chromosome or chromosome fragment, with a cell having a high efficiency for homologous recombination; (b) marking the desired site of insertion of the foreign chromosome using a targeting vector; and (c) inducing deletion or transportation at the marked site. Transport animals produced by the method are useful to provide disease models and knockout animals, and in the production of human proteins, particularly human antibodies. This sequence is used in the method of the invention
                                                                                                                                                                                                                             Producing a cell containing modified foreign chromosomes, useful for the generation of transgenic animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                        Ishida I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 4 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                        Tomizuka K, Yoshida H, Hanaoka K, Oshimura M,
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(KERJ ) FORSCHUNGSZENTRUM JUELICH GMBH.
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                                                                                                                                                                                                                                                                         Example 9; Page 68; 316pp; Japanese
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                                                                                                                                                                                                   WPI; 2000-246479/21.
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                                                                                                                            (KIRI ) KIRIN BEER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-NOV-2000
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                                                                      23-AUG-1999;
                                                                                                21-AUG-1998;
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                                         02-MAR-2000
                                                                                                                                                                       Kuroiwa Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 324
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Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels

3.1%;

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Recombinant Corynebacterium DNA useful for production of pantothenic acid vitamin, comprises panB, panC or ilvD genes encoding enzymes.

Example 1; Page 6; 27pp; German.

This invention describes novel recombinant Corynebacterium DNA (I), present in microrganisms of the Corynebacterium genus and comprising at least one of the panB (ketopantohydroxymethyltransferase), panC (pantothenicacidsyntherase), especially the panBC operon, and/or ilvD (dihydroxyaciddehydratase) genes. (I) is useful for the preparation of pantothenic acid a vitamin which has applications including cosmetics, medicine and human and animal nutrition. The new preparation method using fermentation techniques produces the required stereo-isoform D form of pantothenic acid. This sequence represents a primer used in the isolation of the Corynebacterium glutamicum panBC operon which is described in the method of the invention

Sequence 20 BP; 5 A; 8 C; 5 G; 2 T; 0 U; 0 Other;

ö Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 3; Indels 0; Mismatches Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative

221 GGTGGCGCCAAATCGGG 238 N GTTGTCGGCCACATCGGG 5

ð g RESULT 325

뗦 AAA09167 standard; DNA; 20 AAA09167,

AAA09167;

(first entry) 10-AUG-2000 Nested primer 2 cloning SSH-generated 36P1A6 gene.

36PlA6, transcription factor, murine EHF homologue, ETS family, cytostatic, cancer, vaccine, tumorigenesis; primer, ss. 

Homo sapiens

WO200020584-A2.

13-APR-2000.

99WO-US022576 12-OCT-1999; 98US-0102744P. 99US-0146447P. 02-OCT-1998; 29-JUL-1999;

UROGENESYS INC. (UROG-) (AFAR/)

(AFAR/) AFAR D E. (HUBE/) HUBERT R S. (MITC/) MITCHELL S C.

Afar DE, Hubert RS, Mitchell SC;

WPI; 2000-303772/26

Novel putative transcription factor gene 36P1A6 for treatment, diagnosis and prevention of prostate, bladder, cervical, ovarian, pancreatic, and colonic cancer.

Example 1; Page 30; 53pp; English.

The human 36P1A6 gene encodes a putative transcription factor based on homology to the murine EHF gene which encodes a transcription factor which is a member of the ETS family. 36P1A6 is expressed in androgendeppendent and androgen-independent LAPC prostate cancer xenografts and in normal prostate at approximately equal levels. The highest expression is

in the prostate and colon. 36PlA6 may be involved in activating tumorpromoting genes or repressing genes that block tumorigenesis. The 36PlA6 polynucleotides and polynpeptides are used for the treatment and diagnosis of cancer, e.g. prostate, bladder, cervical, ovarian, pancreatic and colonic cancer (all claimed). Anti-16PlA6 antibodies may be used for burifying 38PlA6 and for isolating 38PlA6 homologues. Antisense oligonucleotides and ribosymes can be used to inhibit the transcription and translation of the 36PlA6 gene (claimed). The 36PlA6 polynucleotides and immunogenic fragments may also be used in cancer vaccines (claimed)

88888888888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ô . 0 Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels 3.1%; 15; Conservative Query Match Best Local Similarity Matches

373 TCCTGGACCGCGACG 390 m recresecededadead

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RESULT 326 AAC64567,

AAC64567 standard; DNA; 20

(first entry) 14-FEB-2001 AAC64567; 

Human prostate specific 30P3C8 nested primer 2 SEQ ID NO:25.

Human, prostate specific gene, 30P3C8, prostate cancer, diagnosis; cytostatic, gene therapy, vaccine, tumour, primer, ss.

Homo sapiens.

WO200061610-A2.

19-OCT-2000.

12-APR-2000; 2000WO-US010218.

12-APR-1999;

(UROG-) UROGENESYS INC.

Raitano AB, Saffran DC Afar DE, Hubert RS, Leong K,

WPI; 2000-619224/59.

30P3CB polypeptide and polynucleotide used for diagnosing, treating and monitoring development of prostate cancer.

Example 1; Page 57; 99pp; English.

The present invention describes human prostate specific protein 30P3C8, which is over-expressed in prostate cancer cells. 30P3C8 has cytostatic cartivity and can be used in vaccines and gene therapy. Methods for detecting the levels of 30P3C8 protein or mRNA in prostate tissue, bone detecting the levels of 30P3C8 protein or mRNA in prostate tissue, bone tissue, lymphatic tissue, serum, blood or semen are used for diagnosing control proteins or cancers in an individual or disregulated cell growth e.g. hyperplasia. The cancers which are detected or diagnosed are of the bladder, pancreas, colon, brain, bone, lung, kidney or prostate by using cell protein, brain, bone, lung, kidney and prostate by using comparers, colon, brain, bone, lung, kidney and prostate of pancreas, colon, brain, bone, lung, kidney and prostate. 30P3C8 colon, brain, bone, lung, cancers expressing colon, brain, bone, lung, cancers. Anti-30P3C8 colon woncolonal antibodies bind to 30P3C8 and disrupt interactions between colons and metastasis and so anti-30P3C8 antibodies may disrupt the homing or and metastasis and so anti-30P3C8 antibodies may disrupt the homing or

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invasion or other cancer promoting activities of 30P3CB. The assays are used for detecting, staging and monitoring prostate cancer. The 30P3CB protein or mRNA are used as additional specific markers for detecting prostate cancer and provide a more specific assay than the serum prostate specific antigen (PSA) assay. The present sequence represents a 30P3CB nested primer, which is used in the exemplification of the present
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G; 2 T; 0 U; 0 Other; Sequence 20 BP; 3 A; 5 C; 10

Gaps ö Ouery Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels rccreeacceceace 390 m recresecescales 373 20 à

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AAC93282 standard; DNA; 20 BP (first entry) 15-FEB-2001 AAC93282; AAC93282 RESULT 

Human STAT3 phosphorothioate antisense oligonucleotide SEQ ID NO:133.

Human, mouse, STAT3; phosphorothioate; antisense oligonucleotide, modulation, signal transducer and activator of transcription; DNA-binding protein; signal transduction; inhibition; apoptosis; inflammatory disease; cancer; antiinflammatory; antirheumatic; cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma; melanoma; lymphoma; diagnosis; ss.

Homo sapiens.

WO200061602-A1.

19-OCT-2000,

06-APR-2000; 2000WO-US009054.

08-APR-1999;

PHARM INC. SISI (-SISI)

Karras JG;

WPI; 2000-619223/59.

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer

Example 12; Page 63; 104pp; English.

The present invention describes an antisense compound (1), 8 to 30 ento-clookes in length, that is targeted to a nucleic acid molecule encoding STAT3 (Signal Transducer and Acrivator of Transcription) and which inhibits the expression of it. (1) has antinflammatory, antirheumatic, cytostatic and immunostimulatory activities. (1) is used for inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are rheumatoid arthritis, cancer of the breast, prostate, brain, had and/or neck, leukaemia, myeloma, melanoma or lymphoma. (I) can also be used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions and for diagnosing the conditions

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(I) can be used alone or with other drugs as an immunostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC932131 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93230 and AAC93232 to AAC93239 represent STAT3 phosphorothioate antisense oligonucleotides, and AAC93300 represents a mismatch control oligonucleotide which are used in example
                                                                                                                                                                                                                                                                                                                                                                                                       modulation, signal transducer and activator of transcription;
DNA-binding protein; signal transduction; inhibition; apoptosis;
inflammatory disease; cancer; antinflammatory; antirheumatic;
cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma;
melanoma; lymphoma; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                  Human STAT3 phosphorothioate antisense oligonucleotide SEQ ID NO:134.
                                                                                                                                                                                                                                                                                                                                                                                              Human; mouse; STAT3; phosphorothioate; antisense oligonucleotide;
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0
                                                                                                                                             Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02;
                                                                                                                                                                      3; Indels
                                                                                                                    Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                136 CCCGCCTGGCGGTGGAGG 153
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                                                                                                                                                                                                                                                                                          В.
                                                                                                                                              3.1%;
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                                                                                            from the present invention
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                                                                                                                                                            Best Local Similarity 83.3
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                 AAC93283;
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                                                                                                                                               Query Match
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06-APR-2000; 2000WO-US009054. WO200061602-A1. Homo sapiens 19-OCT-2000.

08-APR-1999;

(ISIS-) ISIS PHARM INC

Karras JG;

WPI; 2000-619223/59.

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer.

Example 12; Page 63; 104pp; English.

The present invention describes an antisense compound (I), 8 to 30 nucleobases in length, that is targeted to a nucleic acid molecule encoding STATA (Signal Transducer and Activator of Transcription) and which inhibits the expression of it. (I) has antiinflammatory, antirheumatic, cytostatic and immunostimulatory activities. (I) is used for inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are rheumatoid arthritis, cancer of the breast, prostate, brain, head and/or neck, leukaemia, myeloma, melanoma or lymphoma. (I) can also be used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions

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and for diagnosing the conditions associated with expression of STAT3.

(I) can be used alone or with other drugs as an immunostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC93231 encodes mouse STAT3 as given in the exemplification of represent invention. AAC93151 to AAC93230 and AAC93232 to AAC93239 encodes the present sharp hosphorochinate antisense oligonucleotides, and AAC93300 represents a mismatch control oligonucleotide which are used in example
                                                                                                                                                                                                                                                                                                                           3.1%;
                                                                                                                                                                                                                          from the present invention
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Sequence 20 BP; 1 A; 7 C; 8 G; 4 T; 0 U; 0 Other;

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0
Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                              ;
0
                                                          136 CCCGCCTGGCGGTGGAGG 153
                                 15; Conservative
                   Best Loc
Matches
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AAC93216 standard; DNA; 20 (first entry) 15-FEB-2001 AAC93216; **AAC93216**/ RESULT 

BP

Human STAT3 phosphorothicate antisense oligonucleotide SEQ ID NO:67.

modulation; signal transducer and activator of transcription;
DNA-binding protein; signal transduction; inhibition; apoptosis;
inflammatory disease; ancer; antiinflammatory; antirheumatic;
cytostatic; immunostimulatory; rheumatoid arthritis; leukaemia; myeloma;
melanoma; lymphoma; diagnosis; ss. Human; mouse; STAT3; phosphorothioate; antisense oligonucleotide;

Homo sapiens

WO200061602-A1.

19-OCT-2000

06-APR-2000; 2000WO-US009054

99US-0028B461 08-APR-1999;

(ISIS-) ISIS PHARM INC.

Karras JG;

WPI; 2000-619223/59

New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis cancer and

Example 2; Page 47; 104pp; English

The present invention describes an antisense compound (I), 8 to 30 nucleobases in length, that is targeted to a nucleic acid molecule encoding STARIS (Signal Transducer and Activator of Transcription) and which inhibits the expression of it. (I) has antiinflammatory, antitheumatic, cytostatic and immunostimulatory activities. (I) is used for inhibiting the expression of STATIS in cells or tissues, treating an animal having a disease or condition associated with STATIS or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis in a cell. Diseases or conditions that are treated are theuraction arthrities, cancer of the breast, prostate, brain, head and/or neck, lenkaemia, myeloma, melanoma or lymphoma. (I) can also be used for diagnostic methods in detecting and determining the role of

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and for diagnosing the conditions associated with expression of STRT3.

(I) can be used alone or with other drugs as an imminostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC9321 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93330 and AAC93232 to AAC93239 represent STAT3 phosphorothinate antisense oligonuclectides, and AAC933300 represents a mismatch control oligonuclectide which are used in example
                                                                                                                                                                                                                                                                                                                                                                                                                                                   modulation; signal transducer and activator of transcription; bnA-binding protein; signal transduction; inhibition; apoptosis; inflammatory disease; cancer; antiinflammatory; antirheumatic; cytostatic; immunostimulatory; rheumatioid arthritis; leukaemia; myeloma; melanoma; lymphoma; diagnosis; ss.
                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                              Human STAT3 phosphorothioate antisense oligonucleotide SEQ ID NO:47.
                                                                                                                                                                                                                                                                                                                                                                                                                                         mouse; STAT3; phosphorothicate; antisense oligonuclectide;
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                                                                                                                                                                           Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
                                                                                                                                                    Sequence 20 BP; 4 A; 7 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                     292 TGGTGAAGGACCTGAGCC 309
                                                                                                                                                                                                                                                                                                                                   ВР.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-APR-2000; 2000WO-US009054.
                                                                                                                                                                                 Query Match
Best Local Similarity 83.3%;
Matches 15; Conservative
                                                                                                                                                                                                                                                      AAC93196 standard; DNA; 20
                                                                                                                                from the present invention
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                                                                                                                                                                                                                                                                                                          RESULT 33
AAC93196
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New antisense compound for inhibiting the expression of signal transducer and activator of transcription 3 (STAT3) in cells or tissues and treating diseases or condition associated with STAT3, such as rheumatoid arthritis and cancer.

(ISIS-) ISIS PHARM INC.

08-APR-1999;

WPI; 2000-619223/59.

Karras JG;

Example 2; Page 47; 104pp; English.

treated The present invention describes an antisense compound (1), 8 to 30 nucleobases in length, that is targeted to a nucleic acid molecule which inhibits the syzesion of it. (1) has antiinflammatory, antirheumatic, cytostatic and immunostimulatory activities. (1) is used for inhibiting the expression of STAT3 in cells or tissues, treating an animal having a disease or condition associated with STAT3 or a human having a disease or condition associated with STAT3 or a human having a disease or condition characterised by a reduction in apoptosis, and inducing apoptosis, in a cell. Diseases or conditions that are treated are theumatoid arthritis, cancer of the breast, prostate, brain, head and/or neck, leukaemia, myeloma, melanoma or lymphoma. (1) can also be

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used for diagnostic methods in detecting and determining the role of STAT3 in various cell functions, physiological processes and conditions and for diagnosing the conditions associated with expression of STAT3.

(I) can be used alone or with other drugs as an immunostimulator. (I) is used in sandwich and colourimetric assays, involving enzyme conjugation and radiolabeling and is used in diagnostic kits. AAC93150 encodes human STAT3 and AAC93231 encodes mouse STAT3 as given in the exemplification of the present invention. AAC93151 to AAC93230 and AAC93232 to AAC93239 represent STAT3 phosphorothicate antiense oligonucleotides, and AAC93300 represents a mismatch control oligonucleotide which are used in example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          from the present invention
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Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels CCCGCCTGGCGGTGGAGG 153 ö 1 cccecrrecrecrecace 18 3.1%; Conservative Similarity Local St. 15; 136 Query Match Matches g ઠે

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RESULT

AAC64486 standard; DNA; 20

BP

AAC64486;

(first entry) 13-FEB-2001 Prostate tumour associated gene 24P4C12 nested primer 2 SEQ ID NO:41.

Human; prostate tumour associated gene; 24P4Cl2; prostate cancer; transmembrane protein; diagnosis; anticancer; cytostatic; vaccine; gene therapy; PCR primer; ss.

Homo sapiens

WO200061746-A1.

19-OCT-2000.

12-APR-2000; 2000WO-US010039.

99US-0128858P. 12-APR-1999;

(UROG-) UROGENESYS INC

Saffran DC; Raitano AB, Leong K, Afar DE, Hubert RS,

WPI; 2000-672681/65.

Novel 24P4C12 polypeptides and polynucleotides, used in the diagnosis and treatment of cancer, especially prostate cancer.

Example 1; Page 65; 137pp; English.

The present invention describes a prostate tumour associated gene, designated 24P4C12, and its encoded protein. 24P4C12 has anticancer and cytostatic activity, and can be used in vaccine production and in gene therapy. A pharmaceutical composition or vaccine comprising 24P4C12 can be used to treat a patient with cancer, especially prostate cancer, the vaccine can also be used to inhibit the development or progression of cancer. The polypeptides and polynucleotides can be used to diagnose cancers, especially prostate cancer. A transgenic animal comprising 24P4C12 can be used for the development and screening of therapeutic reagents. The polypeptide is a transmembrane protein which is expressed specifically in prostate cancer, allowing the development of more specific anticancer therapies and diagnostic assays

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides the protein and coding sequences of human cancer related protein 20P2H8. The gene, which is found at chromosome 15q32-23, is upregulated in cancers such as that of the prostate, bladder, colon and pancreas. The sequences can be used to diagnose and treat these cancers, and to vaccinate against them. The present sequence is a PCR primer for the coding sequence of the invention
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                          Gaps
                                                                                                                                                                                                                                    Human; cancer related protein 20P2H8; vaccine; chromosome 15q32-23; prostate cancer; bladder cancer; colon cancer; pancreatic cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20P2H8 polynucleotides and polypeptides useful for diagnosing and treating cancer, and for screening for screening for modulating
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 Length 20;
                          Indels
                                                                                                                                                                                                            Human cancer related protein 20P2H8 cDNA PCR primer #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Hubert RS, Mitchell SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene fragment amplifying NP2 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 64; 111pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCCTGGACCGCGACGACG 390
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  3.1%;
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                             Conservative
                                                                                                                                       AAF85709 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Raitano AB,
                                                                                                                                                                                                                                                                                                                                                                                                               (UROG-) UROGENESYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-308645/32.
              Local Similarity
les 15; Conserv
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Best Local S
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                                                                                                                                                                 AAF85709;
     Query Match
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AAD06232/c
                             Matches
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The present invention relates to methods and compositions for the diagnosis and therapy of prostate cancer which utilise human SGP28 (Specific granule protein 28) gene and proteins. The method involves detecting cancers, particularly of prostate and colon, from coverexpression of SGP28 protein. The expression of SGP28, which is an extracted to the prostate and ovary, and is markedly up-regulated in prostate tumours. SGP28 sequence is used for diagnosis (including in vivo imaging), staging, monitoring and prognosis of prostatic and colon cancer, and for assisting selection of therapy. Also SGP28 expressing cancers can be treated by administering a composition or vaccine that contains a vector expressing an antibody specific for SGP28 protein, uncleic acid encoding SGP28 protein or its composition or vaccine that contains a vector expressing an antibody specific for SGP28 protein or therapeutic agent. SGP28 gene product of therapy contingers probed to toxin or therapeutic agent. SGP28 gene product of a also used as source of therapeutic antisense or ribozyme agents, polypeptides encoded by SGP28 gene and SGP28 agent product or is also used as source of therapeutic antisenses or ribozyme agents. SGP28 peptides and for isolating related sequences. SGP28 protein and its caraments are used to raise specific antibodies (Ab) and to identify specific binding agents (potentially useful as therapeutic and diagnostic agents) and also potential anticancer agents. The present sequence is a consetted primer 2 (NP2) used to amplify gene fragments resulting from SSH (suppression subtractive hybridisation) reaction. This sequence is used in the SSH isolation of cDNA fragment of human SGP28 gene
Human, specific granule protein 28, SGP28; therapy; PCR primer; prostate; colon; cancer; prognosis; vaccine; anticancer; SSH; suppression subtractive hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting cancers, particularly of prostate and colon, from overexpression of SGP28 protein, also methods for treating these cancers e.g. by vaccination with the protein.
                                                                                                                                                                                                                                                                                                                                                                     Afar DEH, Mitchell SC, Faris M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 59; 102pp; English
                                                                                                                                                                                                                                   27-OCT-2000; 2000WO-US029607
                                                                                                                                                                                                                                                                                 99US-0162610P
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                                                                                                                                                                                                                                                                                                                                                                         Raitano AB,
                                                                                                                                                                                                                                                                                                                           (UROG-) UROGENESYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-308685/32.
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                                                                                                                                           W0200131343-A2
                                                                                                                                                                                                                                                                                                                                                                         Hubert RS, R
Jakobovits A;
                                                                                                                                                                                                                                                                               28-OCT-1999;
                                                                                               Homo sapiens.
                                                                                                                                                                                    03-MAY-2001
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3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
                                                          373 TCCTGGACCGCGACGACG 390
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                                                                                                                                                          AAH02352 standard; DNA; 20 BP
                                                                               20 TCCTCGGCCGCGACCACG
                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                12-JUN-2001
                                                                                                                                                                                   AAH02352;
                                                                                                                             RESULT 334
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Human AKAP10 coding sequence PCR primer SEQ ID NO: 49.

CXXXXXXXX

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Producing a database for identifying polymorphic genetic markers, comprises obtaining data relating to members of a healthy population and entering the information into a database.
                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides a database of human samples obtained from healthy individuals which can be used to identify polymorphic genetic markers. Data obtained for the database can be used to sort the samples by parameters such as age, sex and ethnicity. This is useful in linking markers with diseases, susceptibility to infection and drug responses. The present primer was used in an assay to demonstrate the uses of the database of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y's disease;
 SNP; human; genetic marker; disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human T-type low voltage activated calcium channel alphalG-c; stress; epilepsy; schizophrenia; depression; sleep disorder; Cushing's disease endocrine disorder; respiratory disorder; peripheral muscle disorder; muscle excitability; fertilisation; contraception; hypertension; neuronal firing regulation; cardiovascular disorder; gene therapy; forensic analysis; epidemiological study; neuroleptic; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18341F PCR primer to generate human calcium channel alphalG-c probe.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                   Ping Y,
                                                                                                                                                                                                                                                                   Koester H, Van Den Boom D,
                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 292; 304pp; English
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Database; polymorphism; SNP; h
drug response; PCR primer; ss.
                                                                                                                                                            13-OCT-1999; 99US-0159176P.
10-UUL-2000; 2000US-0217251P.
10-UUL-2000; 2000US-0217658P.
19-SEP-2000; 2000US-00663968.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-OCT-2000; 2000WO-US027761
                                                                                                                                 13-OCT-2000; 2000WO-US028413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              63 TCTCTGCACTACGAGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAD04754 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 83.3
nes 15; Conservative
                                                                                                                                                                                                                                       (SEQU-) SEQUENOM INC
                                                                                                                                                                                                                                                                                                              WPI; 2001-273865/28.
                                                                                                                                                                                                                                                                                   Jurinke C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200130844-A1
                                                                          WO200127857-A2
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                                             Homo sapiens.
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                                                                                                      19-APR-2001.
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                                                                                                                                                                                                                                                                     Braun A,
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                                                                                                                                                                                                                                                                                     Chiu N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 335
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Matches
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WPI; 2001-308646/32.

Erlander MG;

Zhu JY,

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The invention relates to isoform of human T-type low voltage activated calcium channel (alpha1G-c) cDNA and protein. Cells transformed with calcium channel DNA to express calcium alpha1G-c channel protein are used calcium channel DNA to express calcium alpha1G-c channel protein are used to identify specific modulators (antagonists or agonists). These condulators are useful as therapeutic agents and are used for treating wide range of calcium alpha1G-c channel-mediated disorders, e.g. stress conductive disorders, respiratory disorder, peripheral muscle disorder, chioringly disease, endocrine disorders, respiratory disorder, peripheral muscle disorder, conductive disorders involving regulation, potentiation of synaptic signals and cardiovascular disorders (e.g. atherosclerosis, cardiac hypertrophy, angina pectoris). Calcium alpha1G-c channel DNA is useful conditionally used as antisense sequences, in gene therapy. Calcium channel continually used as antisense sequences, in gene therapy. Calcium channel continually used as antisense sequences, in gene therapy. Calcium channel continually used as antisense sequences, in gene therapy. Calcium channel continually used as antisense sequences, in gene therapy. Calcium channel continually used as antisense sequences, in gene therapy. Calcium channel continually and passays. The present sequence is 18341P PCR primer used for generating calcium alpha1G-c probe which is used for screening human calcium alpha1G-c channel sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, 36P6D5 protein, secreted tumour antigen, therapy, cancer; kidney, bladder, ovary, breast, pancreas, colon, lung, vaccine, cytostatic, SSH; suppression subtractive hybridisation, PCR primer, 88.
                                                                                                                                                    New nucleic acid encoding human calcium channel protein, useful for identifying specific modulators and potential pharmaceuticals for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human 36P6D5 gene fragment amplifying primer NP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;
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Local Similarity 83.3%; Pred. No. 3.8e+02;
les 15; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                       Example 2; Page 45; 115pp; English.
                                                                            Pyati J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 ACTGCCAGTGGCCGAGGG 20
                                        (ORTH ) ORTHO-MCNEIL PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD04811 standard; DNA; 20 BP
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  99US-00426998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 ACTGCGGGTGACCGAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                   treating e.g. epilepsy.
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                                                                              Galindo JE,
                                                                                                                  WPI; 2001-300486/31.
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Mitchell SC;
26-OCT-1999;
                                                                              Dubin AE,
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The present invention relates to a gene and its encoded secreted tumour antigen, termed 36P6D5. These sequences are used for the diagnosis and treatment of various cancers which express 36P6D5, such as cancers of the kidney, bladder, ovary, breast, pancreas, colon and lungs. In normal individuals 36P6D5 protein, is predominantly expressed in pancreas, with comprising immunogenic protein of 36P6D5 is useful for inhibiting the comprising 36P6D5 protein of useful for inhibiting the comprising 36P6D5 protein is useful for diagnosis and/or progness of comprising 36P6D5 genes and/or translation of the 36P6D5 transcripts, and as therappeutic agents. The present sequence is a nested primer (NP)2 and as therappeutic agents. The present sequence is a nested primer (NP)2 used to amplify gene fragments resulting from SSH (suppression the SSH sublation of cDNA fragment of human 36P6D5 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; PC-LECTIN; C-type lectin; transmembrane antigen; normal testis;
layllin homologue; prostate cancer antigen; overexpression;
androgen-dependent prostate cancer; diagnosis; prognosis; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New PC-LECTIN polynuclectide encoding a transmembrane antigen over expressed in human prostate cancer, useful for the prognosis, diagnosis and treatment of prostate cancer.
                                           Detecting presence of cancer expressing 36P6D5 protein in individual by comparing protein level in test sample to normal sample, where elevated level of protein in test sample indicates presence of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer NP2, SEQ ID NO:18, used in human PC-LECTIN cDNA isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                    Example 1; Page 70; 113pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           012/c
AAF76012 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 rctrcdgccgcghcacg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-211222/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
tes 15; Conserv
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Matches
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Gaps

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Length 20; 3; Indels Hubert RS;

Afar DEH,

Faris M,

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contigen, PC-LECTIN (AAB73309) and cDNA encoding it (AAF76004). The expression of the human PC-LECTIN gene is normally restricted to the expression of the human PC-LECTIN gene is normally restricted to the captures of higher in androgen-dependent prostate tumours compared with androgen-independent prostate tumours, and expression is therefore likely cobe dependent on the presence of androgen. Human PC-LECTIN therefore represents a diagnostic and therapeutic target for prostate cancer. Fundancer, correspondent prostate cancer. Human PC-LECTIN exhibits compared to be the human crthologue of layilin, as diverges significantly in a key functional domain proposed for the layilin complexity by the hamman PC-LECTIN or immunogenic portion thereof, a vector protein. Human PC-LECTIN or immunogenic portion thereof, a vector coloning PC-LECTIN or immunogenic portion thereof, a vector coloning PC-LECTIN or immunogenic portion thereof, a vector coloning PC-LECTIN antibodent prostate cancer, but also breast, bladder, lung, bone, particularly prostate cancer, but also breast, bladder, lung, bone, colon, pancreatic, testicular, certical or ovarian cancers that express prostate cancer and other PC-LECTIN expressing cancers. PC-LECTIN proteins are also useful for diagnosing the presence colon, pancreatic, testicular, certical and orlectiones are useful in the transmit (e.g., antisense therapy), diagnosis and/or prognosis of cancer and other PC-LECTIN expressing cancers. PC-LECTIN conclosing and proteins may additionally be used in drug discovery to identify molecules that modulate PC-LECTIN function or expression. The present sequence represents a PCR primer used in the isolation of human property and property of program of the program
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8888888888888888888888888888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps .. 0 Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels ch 3.1%; 1 Similarity 83.3%; 15; Conservative Query Match Best Local Similarity Matches 15; Conserv

373 TCCTGGACCGCGACGACG 390 m rctrceccecada 20 ð 셤

RESULT 338 AAD01985/c

AAD01985 standard; DNA; 20 AAD01985; 

BP

(first entry) 26-MAR-2001

ICV 12 oligonucleotide to construct pMOG845 plasmid.

TPP; bipartite enzyme; trehalose phosphate synthase; trehalose; trehalose phosphate phosphatase; trehalase; transgenic plant; stress resistance; cold; drought; natural flavour; stabiliser; forced water extraction; freeze drying; nutritional value; ss.

Jnidentified.

AU200048921-A

26-OCT-2000

31-JUL-2000; 2000AU-00048921.

97AU-00010085. 09-JAN-1997;

(MOGE-) MOGEN INT NV

Krutwagen RWHH, Voogd E; Goddijn OJM, Verwoerd TC,

WPI; 2001-007580/02.

Chimeric gene encoding bipartite trehalose synthesis enzyme, useful for producing transgenic plants with increased trehalose content.

Disclosure; Page 16; 59pp; English

The present invention relates to a chimeric gene comprising a potato patatin promoter and proteinase inhibitor II terminator (Potrbil), encoding bipartite trehalose synthesising enzyme and method for production of trehalose synthesising enzyme and method for production of trehalose and increasing the level of trehalose cremulation in transgenic plants by inhibiting the degradation of trehalose by trehalose by trehalose phosphate phosphatese (TPS) and trehalose phosphate phosphatese (TPS) and trehalose phosphate phosphatese (TPP) activities, enhances the production of trehalose as it enhalose (TPP) activities, entance to pathway from UPP-glucose and glucose-6-phosphate into trehalose at one and the same site. Plants that contain chimeric gene have improved at one and the same site. Plants that contain chimeric gene have improved at one and the same site. Plants that contain chimeric gene have improved and shelf-life. Trehalose is used for forced water extraction, e.g. in (freeze) drying, particularly where applied to foods, resulting in retention of natural flavours and nutritional value and allowing rapid reconstitution, also as e.g. a stabiliser for vaccines, enzymes, membranes and nucleic acids and it forms a stable, chemically inert construction of pMOG845 plasmid

Seguence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels Conservative Similarity 15; Query Match Local Matches Best

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RESULT 339 AAF83890/

. 0

BP 890/c AAF83890 standard; DNA; 20 AAF83890;

(first entry) 06-AUG-2001 Nested primer (NP)2 used in human PHOR-1 cDNA isolation.

G-protein-coupled receptor; prostate; cancer; PHOR-1; kidney; uterine; cervical; stomach; rectal; cytostatic; vaccine; cell function regulator; human; prostate homologue of olfactory receptor-1; PCR primer; ss.

Homo sapiens.

WO200125434-A1 12-APR-2001 

05-OCT-2000; 2000WO-US027543

99US-0157902P. 05-OCT-1999;

(UROG-) UROGENESYS INC

Jakobovits A, Faris M, Hubert Raitano AB, Afar DEH, J Mitchell SC, Saffran DC;

WPI; 2001-367230/38.

Novel gene designated PHOR-1, a G-protein-coupled receptor up-regulated in prostate cancer, useful as diagnostic marker and therapeutic target for cancers of prostate, kidney, uterus.

Example 1; Page 59; 139pp; English.

The invention relates to a novel G-protein-coupled receptor up-regulated in prostate cancer, termed PHOR-1. The encoding cDNA is contained in plasmid designated pl01P3A11 deposited with ATCC as Accession No.PTA-312. PHOR-1 polypeptides and polymucleotides are useful for diagnosing the presence of cancer, especially prostate, kidney, uterine, cervical,

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(ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                              AAD12168;
                                                                                                                                                                                                                                                                                             RESULT 341
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stomach or rectal cancer by determining and comparing the level of the protein or mRNA expression in teet and normal tissue samples. Pharmaceutical compositions comprising PHOR-1 is useful for treating cancer. PHOR-1 proteins are useful for identifying ligands and other agents and cellular constituents that binds to PHOR-1 gene product and for generating antibodies which are useful in diagnostic, prognostic and imaging methodologies and for the treatment of prostate cancer. Cell innes expressing PHOR-1 are useful for identifying protein-protein interactions mediated by PHOR-1. The present sequence represents a primer used in isolation of the PHOR-1 (prostate homologue of olfactory receptor
                                                                                                                                                                                                                                                                                           Human, PTP1B; protein phosphatase 1B inhibitor; antisense; gene therapy; infection; inflammation; tumour; prophylaxis; phosphorothioate; ss.
                                                                                                                                           Gaps
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                                                                                                                          Length 20;
                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                            Human PTP1B antisense oligonucleotide (ISIS# 107769).
                                                                                                       Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                  note = "Phosphorothioate backbone"
                                                                                                                       Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
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/*tag= b
/mod base= OTHER
/note= "Methoxyethyl residues"
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8. .20
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/mod_base= OTHER
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14. .16
/*tag= f
/mod_base= m5c
16. .20
/*tag= c
                                                                                                                                                            373 TCCTGGACCGCGACGACG 390
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/mod_base= m5c
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                                                                                                                                                                                                                       AAD11960 standard; DNA; 20 BP.
                                                                                                                         3.1%;
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                                                                                                                     Query Match
Best Local Similarity 83.3°
Matches 15; Conservative
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/mod_ba
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                                                                                                                                                                                                                                                                                                                       Homo sapiens,
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                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                         AAD11960;
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                                                                                                                                                                                                      RESULT 34
AAD11960
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                                                                                    New antisense compounds capable of modulating expression of human protein phosphatase 1B, useful for diagnosis, prophylaxis and treatment of diseases associated with expression of protein phosphatase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rat; PTP1B; protein phosphatase 1B inhibitor; antisense; gene therapy; infection; inflammation; tumour; prophylaxis; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rat PTP1B antisense oligonucleotide (ISIS# 111615).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;
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'note= "Phosphorothioate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                   Example 15; Col 42; 71pp; English.
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/mod_base= OTHER
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/mod_base= OTHER
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16..20
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Cowsert LM, Wyatt J;
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                                           WPI; 2001-432181/46.
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modified_base
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The invention is directed to antisense compounds, particularly oligonuclectides which are targetted to a DNA encoding protein phosphatase 1B (PPTB) to modulate its expression. The antisense expension are useful for diagnosis, prophylaxis and treatment of diseases associated with the expression of PTPIB, to prevent or delay infection, inflammation and tumour formation and as a research reagent. The PTPIB DNA is useful in gene therapy. The present sequence is an antisense oligonuclectide with a phosphorchicate backbone. This oligo is targetted to rat PTPIB to inhibit its expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zebrafish; morphogenic signal; neuron; hedgehog gene; embryonic patterning; cell culture; cell differentiation; ischaemia; cell proliferative disorder; intracerebral grafting; Huntington's chorea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compounds capable of modulating expression of human phosphatase 1B, useful for diagnosis, prophylaxis and treatment of diseases associated with expression of protein phosphatase.
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                                                                                                                                                                                                                                             note = "Methoxyethyl residues"
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/mod_base= m5c
16..20
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                                   /*tag= e
/mod_base= m5c
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/mod_base= m5c
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/mod base= m5c
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ID AAD10165 standard; DNA; 20 BP.
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Best Local Similarity 83.3
Matches 15; Conservative
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12-SEP-2001
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EXXXLTXEXEX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention is directed to antisense compounds, particularly oligomucleotides which are targetted to a DNA encoding protein phosphatase 1B (PTP1B) to modulate its expression. The antisense compounds are useful for diagnosis, prophylaxis and treatment of diseases associated with the expression of PTP1B, to prevent or delay infection, inflammation and tumour formation and as a research reagent. The PTP1B DNA is useful in gene therapy. The present sequence is an antisense oligomucleotide with a phosphorothicate backbone. This oligo is targetted to rat PTP1B to inhibit its expression
                                                                                                                                                                                                                                                                                                                                                         New antisense compounds capable of modulating expression of human protein phosphatase 1B, useful for diagnosis, prophylaxis and treatment of diseases associated with expression of protein phosphatase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rat, PTP1B; protein phosphatase 1B inhibitor; antisense; gene therapy; infection; inflammation; tumour; prophylaxis; phosphorothioate; ss.
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/note= "Methoxyethyl residues"
16. .1
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/mod_base= m5c
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                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
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Query Match Best Local Matches 1

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protein

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Gaps

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Synthetic

AAD12156;

RESULT 342 AAD121

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Key
modified_base
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14-DEC-1994;
04-MAY-1995;
05-JUN-1995;
05-JUN-1995;
                    modified_base
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                                   modified base
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                                                  modified base
                                                           US6261786-B1
                                                                    02-JUL-1996;
       Danio rerio
                                                                17-JUL-2001
                                                                                               Marigo V,
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Sequence 20 BP; 3 A; 6 C; 2 G; 1 T; 0 U; 8 Other;
    on 11-SEP-2003 to standardise OS field)
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Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to assay for screening compounds that potentiate or inhibit binding of hedgehog polypeptide to naturally occurring patched receptor. The hedgehog proteans comprise morphogenic signals produced by embryonic patterning centres, and are involved in the formation and maintenance of ordered spatial arrangements of formation and maintenance of ordered spatial arrangements of different discussions in the proteins can be used to generate and/or maintain an array of different vertebrate tissues both in vitro and in vivo. The invention also relates to a method for modulating growth, differentiation or survival of a mammalian cell (e.g. neuron, testicular cell) responsive to hedgehog agonists and antagonists can be used in cell culture techniques to enhance survival and maintenance of neurons and various vertebrate organogenic pathways. The hedgehog gene is useful in determining whether a patient is at the risk of disorder characterised by commend cell prolliferation or aberrant control of differentiation. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Screening compounds that potentiate or inhibit binding of hedgehog polypeptide to naturally occurring patched receptor, comprises contacting polypeptide with receptor and test compound, and detecting change in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hedgehog proteins or mimetics can be used to induce foetal neurons especially neuronal stem cells in intracerebral grafting. The protein or its mimetic can be used in the treatment of neurological conditions e.g. injury to nervous system, ischaemia resulting from stroke, Alzheimer's disease, Parkinson's disease, Huntington's chorea, amyotrophic lateral sclerosis (ALS) and multiple sclerosis. The present sequence is a degenerate PCR primer used to amplify Zebrafish hedgehog gene. (Updated
neurological disorder, Alzheimer's disease; Parkinson's disease;
amyotrophic lateral sclerosis; ALS; multiple sclerosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcmahon AP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD. (HARD ) HARVARD COLLEGE.
                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; Col 91; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ingham PW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93US-00176427.
94US-00356060.
95US-00435093.
95US-00460900.
95US-00462386.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention is related to the coding sequence and protein fragments of a human catenin-binding zinc finger protein. The coding sequence was isolated from a human kidney cDNA library, but is expressed in most human tissue. The sequences provided by the invention can be used in the diagnosis and treatment of cancer and neurological disorders, and in drug screening to identify compounds capable of the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid or its fragments, useful for diagnosing and treating cancer and neurological disorders, corresponds to a catenin-binding protein in signal transduction and gene regulatory pathways.
                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human catenin-binding zinc finger protein PCR primer FVR359R.
3.1%; Score 13.2; DB 1; Length 20; larity 60.0%; Pred. No. 3.8e+02; Conservative 2; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Van Roy F, Vanlandschoot A, Janssens B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 17; 71pp; English
                                                                                                                  133 TGGCCCGCCTGGCGGTGGAG 152
                                                                                                                                                                    20 INGCHMGNYTNGCNGTNGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              286 CCAAGCTGGTGAAGGACC 303
                                                                                                                                                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99EP-00201543,
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Query Match
Best Local Similarity 83.35,
Best Local 15, Conservative
                                                                                                                                                                                                                                                                                                         AAC88711 standard; DNA; 20
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Length 20;

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14-DEC-1994;
                                                                                     US6271363-B1
                                                                                              20-OCT-1997;
                                                                                                           35-JUN-1995;
                                                                                                        34-MAY-1995;
                 Danio rerio.
                                                                                         37-AUG-2001
              Synthetic
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The nucleic acid sequences represent the 83P5G4 gene and the primers and adaptors used to amplify 83P5G4 DNA. 83P5G4 exhibits prostate specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, testis, bladder, kidney, brain, bone, cervix, uterus, ovary, breast, pancreas, stomach, rectum, clon and lung. The 83P5G4 polymuleoide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an single chain monoclonal antibody, that immunospecifically binds to an paying the 83P5G4 related protein, and a ribozyme capable of cleaving a polymucleotide having the 83P5G4 coding sequence, are both useful in the present of a composition for treating a patient with a cancer that expresses 83P5G4. The sequences can be used in diagnostic methods to monitor the level of 83P5G4 gene products in serum, blood, usine and cut issue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           An isolated 83P5G4-related protein useful as a diagnostic and/or therapeutic agent in multiple cancers such as prostate, bladder and bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tumour; kidney; brain; bone; ovary; breast; pancreas; uterus; colon;
lung; cytostatic; gene therapy; antibody therapy; ribozyme; liver;
single chain monoclonal antibody; serum; blood; urine; bladder; cervix;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             83P5G4; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels ...
                                                       6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                             Human prostate-related gene 83P5G4 cDNA nested primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hubert RS, Afar DEH, Challita-Eid PM, Faris M,
Mitchell SC, Jakobovits A;
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       3.1%; Score 13.2; DB 1;
60.0%; Pred. No. 3.8e+02;
tive 2; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rectum; stomach; human; chromosome 1q31-q32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 55; 112pp; English.
                                                                                                  133 rescencecriescesresas 152
                                                                                                                              20 TNGCNMGNYTNGCNGTNGAG 1
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                                                                                                                                                                                                                                                                    BP.
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                                                                                                                                                                                                                                           AAH99163/c
ID AAH99163 standard; DNA; 20
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Query Match
Best Local Similarity 60.0°
Matches 12, Conservative
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nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-514669/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200159115-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                               04-DEC-2001
                                                                                                                                                                                                                                                                                                                  AAH99163;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to nucleic acids encoding hedgehog proteins selected from sonic hedgehog (Shh), indian hedgehog (Ihh), desert hedgehog (Dhh) polypepides. The hedgehog genes are involved in the formation of ordered spatial arrangements of differentiated tissue in vertebrates. The nucleic acid sequences are useful for producing hedgehog proteins, used for promoting differentiation of, or survival of differentiation of neuronal cells, and for promoting proliferation, survival or differentiation of mesenchymal, endodermal or ectodermal tissue, particularly chondrocytes, or testicular germ line cells. Sequences AAH76125-126 represent PCR primers for amplifying a zebrafish Shh genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acid encoding a hedgehog polypeptide, used to produce the polypeptide, which is used to promote proliferation, survival, and/or differentiation of neuronal and mesodermal tissue.
                                                                    Hedgehog protein; sonic hedgehog; Shh; indian hedgehog; Ihh; Dhh; desert hedgehog; cell differentiation; zebrafish; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 6 C; 2 G; 1 T; 0 U; 8 Other;
                      Zebrafish Shh DNA amplifying primer hh 3.3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HARD ) HARVARD COLLEGE.
(IMCR ) IMPERIAL CANCER RES TECHNOLOGY LTD.
                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                        /*tag= b
/mod_base= i
/note= "inosine"
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/mod_base= 1
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94US-00356060.
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/mod_base= i
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95US-00462386
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'note= "Phosphorothioate backbone"

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base= OTHER

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/note= "Methoxyethyl residues"

/\*tag= g /mod\_base= m5c

99US-00467082

Cowsert LM;

base= OTHER

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/\*tag=

16. .20

'\*tag= e note= "Central gap region"

/\*tag= d /mod\_base= m5c

/\*tag= c /mod\_base= m5c

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Antisense oligonucleotides for inhibiting the expression of the human protein kinase A catalytic subunit C-alpha, particularly useful for preventing, delaying or treating infection, inflammation or tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Col 45; 35pp; English.
                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-407321/43.
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                                                                                                                                        misc_feature
                                                                                                                                                                                                                                                                      19-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                    Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                         formation.
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                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to antisense compounds up to 30 nucleobases in length targeted to a E2F transcription factor 1 The invention is useful for inhibiting the expression of E2F transcription factor 1 in cells or tissues. The antisense oligomucleotides may also be used as a research agent and to prevent infection, inflammation or tumours
                                                                                                                    Antisense; E2F transcription factor 1; human; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                            Antisense compound capable of inhibiting the expression of E2F transcription factor 1, useful for preventing or delaying infection, inflammation or tumor formation.
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                                                                                                 Human E2F transcription factor 1 antisense oligonucleotide #71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 3 A; 9 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                       Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                     Example 15; Col 43; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            297 AAGGACCTGAGCCCCGGG 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                m
                                      AAF91365 standard; DNA; 20 BP
                                                                                                                                                                                                           02-MAR-2000; 2000US-00517584.
                                                                                                                                                                                                                              02-MAR-2000; 2000US-00517584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAGGAACTGAGGCCTGGG
                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 83.3
                                                                                                                                                                                                                                                                      Popoff I, Brown-Driver
                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                         WPI; 2001-190981/19.
                                                                                                                                                  Homo sapiens.
                                                                                                                                                                    US6187587-B1
                                                                            04-MAY-2001
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                                                        AAF91365;
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The invention is directed to antisense compounds, particularly oligonucleotides which are targetted to a DNA encoding human protein kinase A (FXA) catalytic subunit C-alpha to modulate (inhibit) its expression. The antisense compounds are useful for diagnostics, therapeutics, prophylaxis and as research reagents or kits. The antisense oligonucleotides are useful for treating human, suspected of having or being prone to a disease or condition associated with the expression of PKA catalytic subunit C-alpha. In particular, the antisense oligonucleotides are useful for preventing, delaying or treating infection, inflammation and tumour formation. They are also useful in antisense therapy. The present sequence is a chimeric antisense oligonucleotide with a phosphorothioate backbone. This oligo is targetted to the cooling region of human PKA catalytic subunit C-alpha to inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 1 A; 8 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  286 CCAAGCTGGTGAAGGACC 303
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Human PKA C-alpha chimeric antisense oligonucleotide (ISIS# 102684).

(first entry)

10-SEP-2001

AAD09658;

658/c AAD09658 standard; DNA; 20 BP.

RESULT 348

AAD09658

Human; protein kinase A; PKA catalytic subunit C-alpha inhibitor; therapy; infection; inflammation; tumour; prophylaxis; antisense; phosphorothioate backbone; chimeric; ss.

Location/Qualifiers

Key

Homo sapiens. Synthetic. Chimeric.

RESULT 349

systemic lupus erythematosus; allograft rejection; ISIS 107231; ss

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                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to human catenin-binding proteins and their corresponding cDNA molecules which functions in signal transduction and gene regulatory pathways. The invention also provides an isolated and/or recombinant nucleic acid or its functional fragment, homologue or derivative, corresponding to a alpha-catenin binding protein. The invention also relates to a novel human zinc finger protein binding with a member of the a-catulin/vinculin family, preferably with a human isoform of alpha N-catenin (neural form). The invention also relates to the field of drug discovery, diagnosis, prognosis and treatment of cancer and neurological disorders. The present sequence is a PCR primer which is used for amplifying human ANC_2HOI CDNA
                                                                                                                                                                                                                                                                                                                                            Novel recombinant nucleic acids useful for diagnosing, prognosing and/or
                                                                                                                                                                                                                                                                                                                                                        treating cancer and neurological disorders, corresponds to a protein binding to alpha-catenin protein and with signal transduction function.
                                                                                                      Human; ANC_2H01 protein; catenin-binding protein; signal transduction; gene regulation; thos finger protein; alphaN-catenin; drug screening; therapy; cancer; neurological disorder; cytostatic; neuroprotective; PCR primer; RACE; rapid amplification of cDNA end; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                  Human ANC_2H01 cDNA amplifying reverse 5' RACE PCR primer, FVR359R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Integrin alpha 4; antisense; very late antigen 4; VLA4; utchind disease; inflammatory disease; rheumatoid arthritis; multiple sclerosis; tumour metastasis; melanoma; asthma; psoriasis; allergy; Grave's disease; Hashimoto's thyroiditis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide for human integrin alpha 4, ISIS 107231.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                          (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                 Van Roy F, Vanlandschoot A, Janssens B;
Example, Page 66; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   286 CCAAGCTGGTGAAGGACC 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ო
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAS10278 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ccaaacrgargaagaacc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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The sequence is an antisense oligonucleotide targetting human integrin 4, a protein involved in autoimmune and inflammatory diseases. The invention relates to antisense inhibitors of integrin alpha 4 which target and inhibit expression of integrin alpha 4. The antisense molecules are useful for inhibiting the expression of integrin alpha 4 in human cells or tissues, treating an animal having a disease or condition associated with autoimmune disease or condition, autoimmune disease or condition; autoimmune disease or condition including rheumatoid arthritis, multiple sclerosis and tumour metastases, melanoma, asthma, psoriasis, allergy, care's disease, Hashimoto's thyroiditis, systemic lugus erythematosus and allograft rejection, and diseases or conditions characterised by leukocyte migration into affected tissues, preferably central nervous system tissues. The antisense molecules are also useful for reducing the levels of VLA-4 and alpha4beta7 integrin in human cells or tissues, and reducing the adherence of cells of a first type e.g., melanoma cells or illicities in the adherence of cells of a first type e.g., melanoma cells or illicities of a second type e.g., melanoma calls or illicities of a second type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities of a first type e.g., melanoma calls or illicities characteria of e.g., melanoma calls or illicities characteria or illic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    οĘ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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                                                                                                                                                                                                             '*tag= a
"mod_base= OTHER
'note= "Other= Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Other= 2' methoxyethoxy residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             methoxyethoxy residues"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 5 C; 12 G; 1 T; 0 U; 0 Other;
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/note= "Other= 2' deoxy residues"
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                                                                                                                           cocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 32; Col 49; 49pp; English.
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/----- "Other= 2"
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/*tag= e
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-450381/48.
                                                                                                                                Key
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Homo sapiens
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                                             Synthetic
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330 GCGGACGACCAGGGCCGG 347

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AAS42202 standard; DNA; 20

AAS42202;

(first entry) 17-DEC-2001 Human prostate-related gene 103P2D6 cDNA nested primer #2.

103P2D6; PCR primer; DNA adaptor; prostate; testis; foetal tissue; ss; tumour; cancer; bone; ovary; breast; pancreas; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; serum; blood; urine; bladder; single chain monoclonal antibody; cervix; human.

Ното варіепв.

WO200162925-A2

30-AUG-2001

26-FEB-2001; 2001WO-US005996.

24-FEB-2000; 2000US-0184558P. 13-JUL-2000; 2000US-0218856P.

(UROG-) UROGENESYS INC

Raitano AB, Afar DEH, Rastegar GS, Mitchell SC, Hubert RS; Challita-Eid PM, Faris M, Jakobovits A;

WPI; 2001-557705/62.

New polynucleotide for treating and diagnosing prostate cancer is the 103P2D6 gene which encodes for 103P2D6-related proteins.

Example 1; Page 55; 132pp; English.

Sequences AAS41293-AAS42208 represent the 103P2D6 gene and the primers and adaptors used to amplify 103P2D6 DNA. 103P2D6 is not expressed in normal adult tissue but is aberrantly expressed in some foreal tissues on and many cancers including tumours of the prostate, testis, bladder, bone, cervix, ovary, breast, pancreas, colon and lung. The 103P2D6 polymucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P2D6 related protein, and a ribozyme capable of cleaving a polymucleotide having the 103P2D6 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P2D6. The sequences can be used in diagnostic methods to monitor the level of 103P2D6 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells 

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

373 TCCTGGACCGCGACGACG 390 ო 20 recressecesceses

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RESULT 352

AAD19416 standard; DNA; 20 BP. AAD19416 ID AAD1 XX

(first entry) 18-DEC-2001 AAD19416; 

Human delta-6-desaturase (hD6D-1) amplifying PCR primer #1.

Delta-6-desaturase gene; D6D; lipid metabolism disorder; atopic eczema; mastalgia; rheumatoid arthritis; Sjogren's syndrome; viral infection; gastrointestinal disorder; post viral fatigue; pre-menstrual syndrome; endometriosis; cystic fibrosis; alcoholism; Alzheimer's syndrome; cardiovascular disease; trohn's disease; congenital liver disease; schizophrenia; diabetic neuropathy; nephropathy; retinopathy; cancer; arterial hypertension; atherosclerosis; chronic inflammatory disorder; autoimmune disorder; hyperholesterolaemia; atopic disorder; hD6D-1; gene therapy; human; PCR primer; ss.

Homo sapiens.

WO200170993-A2

27-SEP-2001.

26-MAR-2001; 2001WO-CA000398

24-MAR-2000; 2000CA-02301158.

(SCOT-) SCOTIA HOLDINGS PLC.

De Antueno RJ; Allen SJ, Ponton A, Smith HL, Winther MD,

WPI; 2001-611507/70.

Nucleic acid encoding delta-6-desaturase gene useful for treating atopic eczema, mastalgia, rheumatoid arthritis, Sjogren's syndrome, gastrointestinal disorders, viral infections and post viral fatigue.

Example 4; Page 69; 164pp; English.

The invention relates to polynuclectides that control delta-6 desaturase genes (D6D) and methods useful for identifying compounds which inhibit or promote the activity of mammalian D6D. Compounds which modulate D6D gene promote the activity of mammalian D6D. Compounds which modulate D6D gene casema, mastalgia, rheumatoid arthritis, Sjogren's syndrome, atopic casema, mastalgia, rheumatoid arthritis, Sjogren's syndrome, endometriosis, vixal infections and post vixal fatigue, premerstrual syndrome, candiovascular disease, Crohn's disease, cancer, Alzheimer's syndrome, cardiovascular disease, Crohn's disease, cancer, compenital liver disease, schizophrenia, diabetes and diabetic compenital invertion are also useful for inhibiting progressive and acute disorders such as arterial hypertension, atherosodierosals, dironic acute disorders and autoimmune disorders, hypertoolesterolaemia and other accopic disorders. D6D genes are useful in gene therapy. The present acquence is a PCR primer used to amplify human delta-6-desaturase (hD6D-

Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels 15; Conservative Query Match Best Local Similarity Matches

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Gaps

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'091/c AAD07091 standard; DNA; 20 RESULT 353 AAD07091/ ID AAD0 XX AC AAD0 XX DT 06-A

(first entry) 06-AUG-2001

AAD07091;

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NP2 primer used in isolation of STEAP cDNA fragment generated from SSH
                  Human, cytostatic, antiproliferative, vaccine, gene therapy, six transmembrane epithelial antigen of the prostate-1, STEAP-1, prostate; colon, bladder; lung, ovarian; pancreatic, PCR primer;
                                                                                                      06-DEC-1999; 99US-00455486.
                                                                                         36-DEC-2000; 2000WO-US033040
                                                                                                                                  Hubert RS,
Jakobovits A;
                                                                                                                    (UROG-) UROGENESYS INC
                                                                                                                                                      WPI; 2001-367804/38.
                                                              WO200140276-A2.
                                                 Homo sapiens.
                                                                                                                                  Afar DEH,
Faris M,
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Raitano AB, Saffran DC, Mitchell SC;

The present sequence is nested primer (NP2) which is used to isolate the human six transmembrane epithelial antigen of the prostate (STEAP) cDNA fragment generated from suppression subtractive hybridisation (SSH3).

STEAP is a member of cell surface serpentine transmembrane antigens.

STEAP gene is used in gene therapy. Inhibiting the development or progression of a cancer (eg. prostate, colon, bladder, lung, ovarian and parceatic) expressing STEAP or inhibiting growth or killing cells expressing STEAP, or inhibiting the prostating a vaccine composition to the patient. Treating a patient with a cancer that expresses STEAP, or inhibiting growth or killing cells expressing STEAP, or inhibiting growth or killing cells expressing sight chains of the monoclonal antibody that comprises the variable domains of the heavy and light chains of the monoclonal antibody that specifically binds to STEAP, such that the vector delivers the single chain monoclonal antibody coding sequence to the cancer cells and the encoded single chain monoclonal antibody is expressed intracellularly New STEAP (six transmembrane epithelial antigen of the prostate) proteins, expressed in human cancers, useful for detecting and treating Example 1; Page 70; 187pp; English.

0; Gaps / Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; nes 15; Conservative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ò 셤 RESULT 354 AAS11672/c ID AAS11672 standard; DNA; 20 AAS11672; 

Prostate and testis-related gene 84P2A9 cDNA nested primer #2. 24-OCT-2001 (first entry)

84P2A9; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; 88; leukaemia; tumour; kidney; brain; bone; skin; ovary; breast; pancreas; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine.

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The mucleic acid sequences represent the 84P2A9 gene and the primers and adaptors used to amplify 84P2A9 DNA. 84P2A9 exhibits prostate and testis specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including leukaemia and tumnours of the prostate, testis, kidney, brain, bone, skin, ovary, breast, pancreas, colon and lung. The 84P2A9 polymucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 84P2A9-related protein, and a ribozyme capable of cleaving a polymucleotide having the 84P2A9 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 84P2A9. The sequences can be used in diagnostic methods to monitor the level of 84P2A9 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                                                                                              New 84P2A9 gene and its encoded protein, useful for diagnosing and treating cancer, e.g. leukemia and cancer of the prostate, testis, kidney, brain or bone, or for eliciting an immune response.
                                                                                                                                                                                                                                            Levin E, Mitchell SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                              Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 71; 149pp; English.
                                                                                                                                                              26-JAN-2000; 2000US-0178560P
                                                                                                                   26-JAN-2001; 2001WO-US002651
                                                                                                                                                                                                                                              Jakobovits A, Afar DEH,
Hubert RS;
                                                                                                                                                                                                        (UROG-) UROGENESYS INC
                                                                                                                                                                                                                                                                                                          WPI; 2001-502631/55.
                                        WO200155391-A2
  Homo sapiens
                                                                                02-AUG-2001
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Gaps 3; Indels 0; Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels Human 158P1F4 gene nested primer (NP)2 SEQ ID NO:736. 373 TCCTGGACCGCGACGACG 390 20 recredecededadeada 3 ABL50419 standard; DNA; 20 BP. 17-JUN-2002 (first entry) ABL50419; RESULT 355 ABL50419, ò

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Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter; 

40200216598-A2

Synthetic.

22-AUG-2001; 2001WO-US026411.

rng.res

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Monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in biological sample.
                                         Challita-Eid PM, Hubert RS,
Faris M, Ge W, Jakobovits A;
      22-AUG-2000; 2000US-0227098P.
10-APR-2001; 2001US-0282739P.
                                                               WPI; 2002-269357/31.
                            (AGEN-) AGENSYS INC
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The present invention describes a method for monitoring 158P1H4 gene products in a biological sample from a patient who has or is suspected of paving cancer. The method comprises determining the status of 158P1H4 gene products in a tissue sample from an individual, comparing the status of the presence of aberrant 158P1H4 gene products in a normal sample, and can the status of 158P1H4 gene products in a normal sample, and contained the presence of aberrant 158P1H4 gene products in the sample. 158P1H4 sequences have cytostatic activity and can be used in vaccine production. 158P1H4 polynucleotides may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic containing the malignant phenotype, in generating and characterising containing the malignant phenotype, in generating and characterising containing the products in normal versus cancerous tissues and so elucidating the malignant phenotype, in generating and characterising cancer vaccines. Antibodies against 158P1H4 are useful in diagnostic and cytotoxic T lymphocyte (CTL) or helper T lymphocyte (HTL) responses, and simmunological reagents for detecting 158P1H4 expressing cells. The as immunological reagents for detecting 158P1H4 expressing cells. The prognostic assays, and imaging methodologies. The 158P1H4 gene has been contended to chromosome 8q22-q23, and the 158P1H4 gene also described in the present invention as represent sequences used in the exemplification of the present invention Example 45; Page 116; 209pp; English.

Ouery Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 373 TCCTGGACCGCGACGACG 390 TCCTCGGCCGCGACCACG 3 ઠે

0; Gaps

ABL50407 standard; DNA; 20 BP (first entry) 17-JUN-2002 ABL50407; RESULT 356 ABL50407/c 셤

Human 158P1H4 gene nested primer (NP)2 SEQ ID NO:724.

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

Homo sapiens. Synthetia. WO200216598-A2

22-AUG-2001; 2001WO-US026411 28-FEB-2002

22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P. 

(AGEN-) AGENSYS INC

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Levin

Raitano AB, Afar DEH,

Raitano AB, Afar DEH, Challita-Eid PM, Hubert RS, F Faris M, Ge W, Jakobovits A;

WPI; 2002-269357/31.

Monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in biological sample.

Example 1; Page 69; 209pp; English.

The present invention describes a method for monitoring 158P1H4 gene products in a biological sample from a patient who has or is suspected of having cancer. The method comprises determining the status of 158P1H4 gene products in a tissue sample from an individual, comparing the status of 158P1H4 gene products in a normal sample, and controlled to the status of 158P1H4 gene products in a normal sample, and controlled to the status of 158P1H4 polynucleotides may be used in wonitoring genetic production. 158P1H4 polynucleotides may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic controlled in ormal versus cancerous tissues and so elucidating the malignant phenotype, in generating and characterising concerting the malignant phenotype, in generating and characterising concerting that bind to 158P1H4 or its particular domain, and for generating cancer vaccines. Antibodies against 158P1H4 are useful in diagnostic and cytotoxic Tlymphocyte (CTL) or helper Tlymphocyte (HTL) responses, and inmaging methodologies. The antibodies are particularly useful in bladder cancer diagnostic and prognostic assays, and imaging methodologies. The also described in correct to chromosome 8922-923, and the 158P1H4 gene also described in the present invention has been located to chromosome 8922-923, and the 158P1H4 gene also described in the present invention has been located to chromosome 8922-923, and the orenomosome 8923. Ablished to chromosome 8922-923, and the 158P1H4 gene also described in the exemplification of the present invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels Query Match
Best Local Similarity 83.3
Matches 15; Conservative

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373 TCCTGGACCGCGACGACG 390 

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AAS96899 standard; DNA; 20 BP. 20 RESULT 357 AAS96899 셤

AAS96899;

Human STAT3 antisense phosphorothioate oligodeoxynucleotide #106. (first entry) 26-FBB-2002

sTAT3; human; signal transducer and activator of transcription; ss; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breast; prostate; head; neck; brain; leukaemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic.

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Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.
                                                                                                                                                                Example 12; Page 18; 21pp; English
                                                                 08-APR-1999; 99US-00288461.
06-APR-2000; 2000WO-US009054.
                                                  11-JAN-2001; 2001US-0075881
                                                                                                                    WPI; 2002-009991/01.
                                                                                       (KARR/) KARRAS J G.
                      US2001029250-A1.
Homo sapiens
                                    11-OCT-2001
        Synthetic.
                                                                                                      Karras JG;
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The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAT) protein, specifically STAT3, where the antisense compounds inhibit the expression of STAT3. The antisense squences are useful for inhibiting the expression of STAT3 in cells or tissues, inducing Fas-mediated approasis in cells, and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAT3. These disorders include inflammatory or autoimmune breast, prostate, brain and head and neck and leukaemias, myelomas, melanomas and lymphomas. Also treatable are human diseases or conditions characterised by a reduction in apoptosis or an insensitivity to apoptotic signals. The sequences of the invention can be used in clinical research, for detecting and determining the role of STAT3 in various cell functions and physiological processes and for diagnosing conditions canced with the expression of STAT3. The sequences represent cDNA encoding human STAT3 and human STAT3 oligonucleotides Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;

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0; Gaps
           3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels
                                                                        136 CCCGCCTGGCGGTGGAGG 153
Query Match
Best Local Similarity bo...
These 15; Conservative
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cccccrrccrccrccacca 19 AAS96900 standard; DNA; 20 BP AAS96900; RESULT 358 AAS96900 g

Human STAT3 antisense phosphorothioate oligodeoxynucleotide #107. 

(first entry)

26-FEB-2002

STAT3; human; signal transducer and activator of transcription; ss; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breast, prostate; head; neck; brain; leukaemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic

Homo sapiens. Synthetic.

Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is taxgeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins. Example 12; Page 18; 21pp; English 11-JAN-2001; 2001US-00758881 08-APR-1999; 99US-00288461. 06-APR-2000; 2000WO-US009054. WPI; 2002-009991/01 (KARR/) KARRAS J G. US2001029250-A1 11-OCT-2001 Karras JG; 

The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAMI) protein, specifically STAMI3, where the antisense compounds inhibit the expression of STAMI3 in cells are useful for characteristic and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAMI3. These disorders include inflammatory or autoimmune of disease, particularly rheumatoid arthritis, cancers, such as those of the breast, prostate, brain and head and neck and leukaemias, myelomas. Or characterised by a reduction in apoptosis or an insensitivity to apoptotic signals. The sequences of the invention can be used in clinical capacity for detecting and determining the role of STAMI3 in various cell functions and physiological processes and for diagnosing conditions associated with the expression of STAMI3. The sequences concerned and for diagnosing conditions associated with the expression of STAMI3. The sequences of the conditions associated with the expression of STAMI3. The sequences of the conditions associated with the expression of STAMI3. The sequences of the conditions are second to be stated to be stated to be associated with the expression of STAMI3. The sequences of the conditions are second to be stated to be stated to be associated with the expression of STAMI3. The sequences represent cDNA encoding human STAMI3 oligonucleotides

0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels Sequence 20 BP; 1 A; 7 C; 8 G; 4 T; 0 U; 0 Other; 136 CCCGCCTGGCGGTGGAGG 153 Query Match
Best Local Similarity 85...
The local Similarity 15...
The local Similarity 15... ઠે

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Human STAT3 antisense phosphorothioate oligodeoxynucleotide #66. 3 cccecrradredredace 20 AAS96833 standard; DNA; 20 BP. 26-FEB-2002 (first entry) AAS96833; RESULT 359 AAS96833/

STAT3; human, signal transducer and activator of transcription; ss; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breast; prostate; head; neck; brain; leukaemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; neck; brain; leukaemia; myeloma; melanoma; US2001029250-A1. Homo sapiens. cytostatic Synthetic. 

11-0CT-2001

Wed Apr 21 12:58:21 2004

08-APR-1999; 99US-00288461. 11-JAN-2001; 2001US-00758881

(KARR/) KARRAS J G.

WPI; 2002-009991/01.

Karras JG;

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The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAMI) specifically STAMI, where the antisense compounds inhibit the expression of STAMI; where the antisense compounds inhibit the expression of STAMI in cells or tissues, inducing Fascindiated apoptosis in cells, and sensitising cells to apoptosis. They are also useful for treating an animal having a disease or condition associated with STAMI. These disorders include inflammatory or autoimmune of disease, particularly rheumatoid arthritis, cancers, such as those of the breast, prostate, brain and head and neck and leukaemias, myelomas. Compared the parterised by a reduction in apoptosis or an insensitivity to apoptotic signals. The sequences of the invention can be used in clinical cappetotic signals. The sequences of the invention can be used in clinical cunctions and physiological processes and for diagnosing conditions associated with the expression of STAMI. The sequences represent cDNA encoding human STAMI and human STAMI oligonuclectides
Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.
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                                  Gaps
                                  ö
3.1%; Score 13.2; DB 1; Length 20;
83.3%; Pred. No. 3.8e+02;
tive 0; Mismatches 3; Indels
    Query Match
Best Local Similarity 83.3'
Matches 15; Conservative
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AAS96813 standard; DNA; 20 BP RESULT 360 AAS96813

AAS96813;

(first entry) 26-FEB-2002 Human STAT3 antisense phosphorothioate oligodeoxynucleotide #46.

STAT3; human; signal transducer and activator of transcription; se; STAT; antisense gene therapy; Fas-mediated apoptosis; inflammatory disease; autoimmune disease; rheumatoid arthritis; cancer; breat; prostate; head; neck; brain; leuksemia; myeloma; melanoma; lymphoma; apoptosis; antiinflammatory; immunosuppressive; antirheumatic; antiarthritic; cytostatic.

Homo sapiens. Synthetic.

US2001029250-A1.

11-OCT-2001.

11-JAN-2001; 2001US-00758881 

08-APR-1999; 99US-00288461. 06-APR-2000; 2000WO-US009054. 

(KARR/) KARRAS J G.

Karras JG;

WPI; 2002-009991/01

Novel antisense compound useful for treating and diagnosing inflammatory diseases and cancers, is targeted to a nucleic acid molecule encoding signal transducer and activator of transcription proteins.

Example 2; Page 13; 21pp; English.

The invention relates to antisense compounds targeted to a nucleic acid molecule encoding a signal transducer and activator of transcription (STAT) by protein, specifically STAT3, where the antisense compounds inhibit the expression of STAT3. The antisense squences are useful for the expression of STAT3 in cells or tissues, inducing Fasconditions also useful for treating an animal having a disease or condition associated with STAT3. These disorders include inflammatory or autoimmune of disease, particularly theumatoid arthritis, cancers, such as those of the breast, prostate, brain and head and neck and leukaemias, myelomas, or conditions and lymphomas. Also treatable are human diseases or conditions or apoptotic signals. The sequences of the invention can be used in clinical conditions and physiological processes and for diagnosing conditions or functions and physiological processes and for diagnosing conditions associated with the expression of STAT3. The sequences represent cDNA encoding human STAT3 and human STAT3 oligonucleotides

Sequence 20 BP; 2 A; 5 C; 9 G; 4 T; 0 U; 0 Other;

0; Gaps 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indel8 Query Match Best Local Similarity 83.3 Matches 15; Conservative

ð g RESULT 361 AAS62190

86 AAS62190 standard; DNA; 20

AAS62190;

(first entry) 29-JAN-2002 Porcine forward PCR primer for bFGF.

metabolic; muscular; MSMF; Pig; muscular steatosis-modulating factor; ss; metabolic; muscular; MS food supplement; obesity; hyperlipidaemia; atherosclerosis; wound healing; tumour; amyotrophic lateral sclerosis; ALS; PCR primer.

Sus scrofa.

WO200179287-A2.

HANGE STANDARD STANDA

25-OCT-2001

12-APR-2001; 2001WO-CA000509.

17-APR-2000; 2000US-0197936P.

MIAC ) CANADA AGRIC & AGRI-FOOD CANADA.

Gariepy C; Pomar C, Palin M,

WPI; 2002-017600/02

Prognosis and diagnosis of muscular steatosis, useful e.g. for selecting animals for breeding, by measuring levels of specific markers, also treating or inducing steatosis.

Example 1; Page 39; 190pp; English.

The invention relates to prognosis or diagnosis of muscular steatosis by measuring the level of a muscular steatosis modulating feator (MSMF) in a human or animal and comparing this with the level in a healthy control. Any difference indicates presence of, or predisposition to, muscular steatosis. The method is particularly used for diagnosis or prognosis of counders in animal breeding. Also (ant) agonists of MSMF can be used to treat, or induce (for increasing the fat content of food) muscular steatosis, in humans and animals. The MSMF markers are also useful in the study of diseases and conditions such as obesity, hyperlipidaemia, archeroscierosis, wound healing, tumours and amyotrophic lateral sclerosis invention from its gene

Sequence 20 BP; 2 A; 7 C; 6 G; 5 T; 0 U; 0 Other;

Gaps ö Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels 0; Mismatches 3.1%; 15; Conservative Query Match Best Local Similarity Matches

237 GGAGGCTGCTTCCCGGGC 254 3 GGAGGCTTCTTCCTGCGC 20 à

RESULT 362 ABA98342,

ABA98342 standard; DNA; 20 29-NOV-2002 (first entry) ABA98342; 

55P4H4; cancer; immune response; ds; PCR primer. Nested primer (NP) 2.

WO200196391-A2. Unidentified.

13-JUN-2001; 2001WO-US019246. 20-DEC-2001.

13-JUN-2000; 2000US-0211454P. (UROG-) UROGENESYS INC.

WPI; 2002-098053/13.

Jakobovits A;

Faris M,

Novel isolated 55P4H4-related protein encoded by a gene over-expressed in multiple cancers, useful as a diagnostic and/or therapeutic agent for cancer, preferably prostate cancer.

Example 1; Page 54; 160pp; English.

This invention relates to an isolated 55P4H4-related protein encoded by a gene that is over-expressed in multiple cancers. The polypeptide is useful for inducing an immune response to an 55P4H4 protein, providing the protein comprises of at least one T cell or B cell epitope. The immune system cell is a B cell which generates antibodies that specifically bind to the protein or is a T cell, preferably a cytotoxic T

cell (CTC) which kills an autologous cell that expresses the 55P4H4
protein, or a helper T cell (HTL) which secretes cytokines that
callitate the cytotoxic activity of a cytotoxic T lymphocyte. A method
is mentioned which is considered useful for monitoring the presence of
cancer in an individual, where the presence of elevated 55P4H4 mRNA or
protein expression in the test sample relative to the normal tissue
sample provides an indication of the presence or status of cancer which
course in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
ocurse in a prostate, kidney, testis, lung, cervix, bone, bladder, brain
conditions associated with disregulated cell growth such as cancer and is
also useful in forenaic analysis of tissues of unknown origin, to treat a
pathological condition characterized by the overaxpression of 55P4H4, for
assessing the status of 55P4H4 gene products in normal versus cancerous
tissue, and to assess the presence of perturbations in specific regions
of the 55P4H4 gene. This sequence represents nested primer (NP) 2 used
during the method highlighted in the examples

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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; Live 0; Mismatches 3; Indels Query Match
Best Local Similarity 83.33
Matches 15, Conservative

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RESULT 363 ABL41837

ABL41837 standard; DNA; 20 BP.

ABL41837;

(first entry) 29-MAY-2002

PCR primer for rat endometriotic protein ENDO-I cDNA.

Rat; endometriotic protein; ENDO-1; glycoprotein; stromal cell; endometriotic tissue; endometriosis; PCR primer; ss.

Rattus sp Synthetic

US2002009718-A1.

24-JAN-2002

98US-00044604. 19-MAR-1998; 

94US-00328451. (TIMM/) TIMMS K L. 25-OCT-1994;

Timms KL;

WPI; 2002-215823/27.

Raitano AB;

Mitchell SC,

Levin E,

Hubert RS, Afar DEH,

Novel purified and isolated glycoprotein designated ENDO-1, useful as marker for diagnosing endometriosis in female patient suspected of having endometriosis,

Example 7; Page 10; 20pp; English.

PCR primers ABL41837-38 were used to amplify a cDNA fragment of rat endometrictic protein ENDO-1. ENDO-1 is a N-acetyl linked glycoprotein, synthesized and secreted specifically by stromal cells of endometrictic tissue origin. Human ENDO-1 has a molecular weight of 40000-55000 as determined by two-dimensional sodium dodecyl sulphate-polyacrylamide gelectrophoresis (SDS-PAGE), and has an isoelectric point of 4.0-5.5. ENDO-1 is useful as a marker for diagnosing endometriosis in a female patient suspected of having endometriosis. Endometriosis in a female patient be diagnosed by obtaining a sample from the patient, and detecting the

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targeted to a nucleic acid encoding protein phosphatase 1B (PTP1B), where
the compound specifically hybridises with and inhibits the expression of
PTP1B (e.g. an antieense oligonucleotide). Also included are (1) a
COMPOUND of an active site on a nucleic acid encoding
COMPOUND of an active site on a nucleic acid encoding
COMPOUND of an active site on a nucleic acid encoding
COMPOUND of an active site on a nucleic acid encoding
COMPOUND or suspected of Paying alsease or condition associated
COMPOUND or suspected of having a disease or condition associated
COMPOUND or an animal comprising administering the compound; (5)
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  presence of ENDO-I in the sample compared to non-endometriosis controls
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense; protein phosphatase 1B; PTP1B; ss; probe; rat; type 2 diabetes; obesity; ovarian cancer; chronic myeloid leukaemia; hyperproliferative disease; antidiabetic; anorectic; cytostatic; blood glucose; gene therapy.
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Pred. No. 3.8e+02;
0; Mismatches 3; Indels
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                                               Seguence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat PTPB1 antisense oligonucleotide ISIS 111603.
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                                                                                                                                    0;
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31-JUL-2000; 2000US-00629644.
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                                                                                  Query Match
Best Local Similarity 83.35
Matches 15; Conservative
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FREIER S M.
MONIA B P.
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                                                                                                                                                                                                                                                                                                                                                                                                            ABK85231;
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                                                                                                                                                                                                                                                                                                                  RESULT 364
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The invention relates to a compound of 8-50 nucleobases in length targeted to a nucleic acid encoding protein phosphatase 1B (FTP1B), where the compound specifically hybridises with and inhibits the expression of PTP1B (e.g. an antisense oligonucleotide). Also included are (1) a compound of 8-50 nucleobase oligonucleotide). Also included are (1) an 8 nucleobase portion of an active site on a nucleic acid encoding PTP1B; (2) inhibiting the expression of FTP1B in cells or tissues comprising contacting the cells or tissues with the compound; treating an animal having or suspected of having a disease or condition associated with PTP1B comprising administering the compound; (4) decreasing blood
                                                                                                                                                                                                                                                                                                                                                                                                            ö
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pTPIB, such as type 2 diabetes, obesity, cancer (especially ovarian cancer, chronic myeloid leukaemia and hyperproliferative diseases in an animal having or suspected of having the disease or condition, and decreasing blood sugar levels or preventing or delaying the onset of an increase in blood glucose levels in an animal. The compound is also used in diagnoserics, therapeutics, prophylaxis, and in research reagents and kits. The present sequence is an antisense compound of the invention targetting rat PTPIB
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense, protein phosphatase 1B; PTP1B; ss; probe; rat; type 2 diabetes; obesity; ovarian cancer; chronic myeloid leukaemia; hyperproliferative disease; antidiabetic; anorectic; cytostatic;
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                                                                                                                                                                                                                                                                                     Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                 Score 13.2; DB 1;
Pred. No. 3.8e+02;
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31-JUL-2000; 2000US-00629644.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   blood glucose; gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK85243 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                             3.14
Query Match
Best Local Similarity 83.33
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COWSERT L M.
WYATT J.
FREIER S M.
MONIA B P.
BUTLER M M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-462914/49.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus norvegicus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2002055479-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK85243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MONI/)
(BUTL/)
(MCKA/)
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(WYAT/)
(FREI/)
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cengar levels in an animal comprising administering the compound; (5)
preventing or delaying the onset of a disease or condition associated
with PPPID in an animal comprising administering the compound; and (6)
preventing or delaying the onset of an increase in blood glucose levels
in an animal comprising administering the compound. The compound is used
to inhibit the expression of PPPIB in cells or tissues, to treat or
prevent or delay the onset of a disease or condition associated with
PPPIB, such as type 2 diabetes, obesity, cancer (especially ovarian
cancer, chronic myeloid leukaemia and hyperprofiferative diseases in an
mimal having or suspected of having the disease or condition, and for
decreasing blood sugar levels or preventing or delaying the onset of an
increase in blood glucose levels in an animal. The compound is also used
in diagnostics, therapeutics, prophylaxis, and in research reagents and
kits. The present sequence is an antisense compound of the invention

Sequence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;

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3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels
                                                                        270 CTGGAGCAGGGGGCACC 287
                                                                                            19 CTGGAGCAGGCCAGGACC 2
3.17
Best Local Similarity 83.33
Matches 15, Conservative
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Gaps

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ABK85035 standard; DNA; 20 BP. 

(first entry) 13-AUG-2002 ABK85035;

Human PTP1B antisense oligonucleotide ISIS 107769.

Antisense; protein phosphatase 1B; PTP1B; ss; probe; human; type 2 diabetes; obesity; ovarian cancer; chronic myeloid leukaemia; hyperproliferative disease; antidiabetic; anorectic; cytostatic; blood glucose; gene therapy.

Homo sapiens.

JS2002055479-A1.

09-MAY-2002.

14-MAY-2001; 2001US-00854883.

18-JAN-2000; 2000US-00487368.

COWSERT L M.

WYATT J. FREIER S M. MONIA B P. BUTLER M M. COWS/)
WYAT/)
FREI/)
MONI/)

MCKAY R.

MCKA/)

Freier SM, Monia BP, Butler MM, Mckay R; Wyatt J, Cowsert LM,

WPI; 2002-462914/49.

Compound for inhibiting the expression of protein phosphatase 1B (PTP1B) and for treating diabetes, cancer, or obesity, comprises an antisense oligonucleotide targeted to nucleic acid encoding PTP1B.

Example 15; Page 23; 133pp; English.

The invention relates to a compound of 8-50 nucleobases in length targeted to a nucleic acid encoding protein phosphatase 1B (PTP1B), where the compound specifically hybridises with and inhibits the expression of

CC compound of 8-50 nucleobases in length which specifically hybridises with an 8 nucleobase portion of an active site on a nucleic acid encoding property (2) inhibiting the expression of PTP1B in cells or tissues comprising contacting the captesion of PTP1B in cells or tissues comprising contacting the calls or tissues with the compound; treating an animal having or suspected of having a disease or condition associated with PTP1B comprising administering the compound; (4) decreasing blood sugar levels in an animal comprising administering the compound; (5) preventing or delaying the onset of a disease or condition associated with PTP1B in animal comprising administering the compound; and (6) preventing or delaying the onset of an increase in blood glucose levels in an animal comprising administering the compound; The compound is used to inhibit the expression of PTP1B in cells or tissues, to treat or prevent or delay the onset of a disease or condition associated with PTP1B, such as type 2 diabetes, obesity, cancer (especially ovarian cancer; chronic myeloid leukaemia and hyperproliferative diseases in an animal having or suspected of having the disease or condition, and for decreasing blood sugar levels or preventing or delaying the onset of an increase in blood glucose levels in an animal. The compound is also used in diagnostics, therapeutics, prophylaxis, and in research reagents and kits. The present sequence is an antisense compound of the invention cut argetting human PTPIB

Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels (

303 CIGAGCCCCGGGGACCGC 320 1 crradccccdadcccccc 18 ઠે 셤

RESULT 367 ABA03609/

멾. ABA03609 standard; DNA; 20

ABA03609;

(first entry) 08-FEB-2002 Nested primer 2 used for human 34P3D7 cDNA isolation.

Human, 34P3D7; cytostatic; vaccine; gene therapy; cancer; human leukocyte antigen; HLA; major histocompatibility complex; MHC; HLA A1; HLA A11; HLA A02; HLA A24; HLA A3; HLA B35; HLA B7; primer; 88.

Homo sapiens.

WO200159110-A2. 

16-AUG-2001.

08-FEB-2001; 2001WO-US004094.

08-FEB-2000; 2000US-0181020P.

(UROG-) UROGENESYS INC.

Challita-Eid PM, Hubert RS, Levin Faris M, Afar DEH, Challi Mitchell SC, Jakobovits A;

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WPI; 2002-025689/03.

New gene designated 34P3D7, encoding a tissue-specific protein highly expressed in prostate cancer, for use as diagnostic and/or therapeutic target for cancers, and for eliciting an immune response.

Example 1; Page 53; 112pp; English.

The invention relates to a polynucleotide, designated 34P3D7, encoding 34P3D7-related protein, comprising a sequence of 2198 nucleotides fully

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Gaps

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defined in the specification. The presence of elevated 3493D7 mRNA or protein expression indicates the presence of cancer occurring in brostate, bladder, kidney, brain, bone, cervical, uterine, ovarian, breast, pancreatic, stomach, colon, rectal leukcoytes, liver, and lung tissue, and in melanocytes. An antibody against the 3493D7-related polynucleotide, or a ribozyme capable of cleaving the 3493D7 polynucleotide, or a ribozyme capable of cleaving the 3493D7 polynucleotide, or a ribozyme capable of cleaving the 3493D7 polynucleotide is useful for inhibiting the development of a cancer expressing 3493D7 in a patient. The present sequence was used in an example demonstrating suppression subtractive hybridisation (SSH)-generated isolation of a cDNA fragment of the 3493D7 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; calreticulin; antisense compound; hyperproliferative disorder; cancer; autoimmune disease; viral infection; cardiovascular disease; antisense therapy; cytostatic; immunosuppressive; virucide; antisense; phosphorothioate backbone; ss.
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                                                                                                                                                                                                                                                                                                                                                                                    Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
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note= "2'methoxyethyl nucleotides"
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                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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/note= "Phosphorothioate backbone"
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/mod_base= m5c
6..20
/*tag= c
/mod_base= OTHER
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/mod_base= m5c
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/mod_base= m5c
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/mod_base= m5c
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/mod_base≈ m5c
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Best Local Similarity
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modified_base
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Synthetic.
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The invention relates to antisense compounds, compositions and methods for modulating the expression of calreticulin. The compositions comprise antisense compounds, particularly antisense oligonuclectides, targetted to nucleic acids encoding calreticulin. The antisense compound is useful for inhibiting the expression of calreticulin in human cells or tissues. It is also useful for treating a human having a disease or condition associated with calreticulin, e.g., hyperproliferative disorder e.g. cancer, autoimmune disease, viral infection or cardiovascular disease, by inhibiting expression of calreticulin. It is useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. It is also used in antisense therapy. The present sequence is an antisense compound arracetted to human calreticulin. This sequence is used to study the antisense inhibition of calreticulin expression-phosphorothioate 2. MOE gapmer oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, 125P5C8; cancer, cytostatic, breast cancer, prostate cancer;
bladder cancer; kidney cancer, colon cancer; ovarian cancer; PCR; primer;
                                                                                                                                                                                                                                                                       Novel antisense compound targeted to nucleic acid encoding calreticulin, useful for treating a human having disease or condition associated with calreticulin e.g. cancer, viral infection, autoimmune disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 83; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            342 GGCCGGCTGCTCTACAGC 359
/*tag= j
/mod_base= m5c
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                                                                                                            30-OCT-2001; 2001WO-US049045
                                                                                                                                            30-OCT-2000; 2000US-00702327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGCAGGCCTCTCTACAGC
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AAL50002 standard; DNA; 20
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Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                             Cowsert LM;
                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                             WPI; 2002-479759/51.
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                                               WO200236743-A2
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                                                                                                                                                                                                           Bennett CF,
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                                                                             10-MAY-2002
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Afar DEH, Raitano AB,

Hubert RS, Jakobovits A;

(AGEN-) AGENSYS INC.

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The present invention relates to compositions comprising a substance that modulates the status of 125P5C8 or a molecule that is modulated by 125P5C8. The status of a cell that expresses 125P5C8 is modulated. The composition is useful for treating cancer, particularly prostate, bladder, kidney, colon, ovary or breast cancer. The 125P5C8 protein and/or a nucleotide sequence encoding the protein is useful for immunising a mammal against cancer. The present sequence is a PCR primer shown in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human/mouse C/EBP phosphorothioate antisense oligonucleotide, SEQ ID:41
                                                                                                                     New composition comprising a substance that modulates the status of 125P5C8 gene or a molecule that is modulated by 125P5C8, useful for treating or preventing cancer that expresses or over expresses 125P5C8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, C/BBP alpha; CCAAT/enhancer-binding protein alpha; CEBPA; transcription factor; tissue development; cellular function; prollferation, differentiation, adipocyte; energy metabolism; chondrogenic; ovulation; follicular development; chordrogenic; ovulation; follicular development; hormonal metabolic regulation; paranlocyte development; cancer; hormonal metabolic regulation; granulocyte development; cancer; tumour formation; infection; infantation; expression inhibition; antisense therapy; quantitative real-time PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                      Query March
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                             Example 1; Page 68; 274pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  373 TCCTGGACCGCGACGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABA02229 standard; DNA; 20 BP.
                                  Faris M, Challita-Eid PM,
Morrison RK, Morrison K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-FEB-2002 (first entry)
                                                                                     WPI; 2002-713510/77.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 370
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20 TCCTCGGCCGCGACCACG 3

1..20 Arags a /mod bass= OTHER /note= "Phosphorothioate linkages"

'mod\_base= OTHER

\*tag= b

16. .20 /\*tag= c

modified base

JS6306655-B1

Location/Qualifiers

Gaps

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cothe human CCAAT/enhancer-binding protein alpha (C/BBP alpha) gene, which human CCAAT/enhancer-binding protein alpha (C/BBP alpha) gene, which inhibit it lits expression. The antisease oligonucleotides were designed to target different regions of the human C/BBP alpha RNA, and were analysed for their effect on C/BBP alpha mRNA levels by quantitative captelled for their effect on C/BBP alpha mRNA levels by quantitative captelled of hybridising to mouse C/BBP alpha mRNA. The C/BBP family of proteins are a family of transcription factors which regulate the expression of wide range of genes that control normal tissue development, callular function, of genes that control normal tissue development, callular function, and service and chones feration and functional differentiation.

C/BBP alpha (also known as CBBPA) is primarily found in tissues involved in energy metabolism which have a capacity to metabolise involved in collecter activity, in the hormonal regulation of adjocyte and chondrogenic differentation, and is also involved in follicular development and ovulation, steroid-induced cell cycle arrest in the liver, in controlling glucose transporter GUTZ promoter activity, in the hormonal regulation of metabolism, and in cycle arrest in the liver, in controlling clucose transporter GUTZ promoter activity, in the hormonal regulation of metabolism, and in cycle arrest promoter activity, and the diagnosis, prevention and treatment of conditions associated with control of conditions associated with control of conditions associated with control of conditions are useful controlled to the conditions of conditions or controlled cont
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                                                                                                                                                                                                                                                                                          New antisense oligonucleotides for modulating the expression of CCAAT/Enhancer-binding proteins alpha, particularly useful for preventing, delaying or treating infection, inflammation or tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3, Indels 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cch 3.1%; Score 13.2; DB 1; Length 20; al Similarity 83.3%; Pred. No. 3.8e+02; 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human cancer-related gene 103P3E8 cDNA nested primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 9 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                         Example 15; Col 42; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 335 CGACCAGGGCCGGCTGCT 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CGGCCAGCGCCAGCTGCT 19
                                                                                                                                                                                            Monia BP, Butler MM, Wyatt J;
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                                             13-JUN-2000; 2000US-00593589.
                                                                                              13-JUN-2000; 2000US-00593589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS95820/c
ID AAS95820 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                               WPI; 2002-040202/05.
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200179557-A2
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23-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS95820;
                                                                                                                                                                                                                                                                                                                                                                              formation
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/note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE
cytosines are 5-methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' MOE cytosines are 5-methylcytosine"
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Wed Apr 21 12:58:21 2004

(UROG-) UROGENESYS INC

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Example 1; Page 55; 128pp; English.
                                                                                                                                                                                                                            AAS99443 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                             24-MAY-2001; 2001WO-US017495
                                                                                                                                                                                                                                                                                                                                       24-MAY-2000; 2000US-0207138P
   12-APR-2001; 2001WO-US012181.
             12-APR-2000; 2000US-0196647P.
                                                                                                                                                                                                                                               12-MAR-2002 (first entry)
                      (UROG-) UROGENESYS INC
                                             WPI; 2002-061976/08.
                                                                                                                                                                                                                                                                                                            WO200190157-A2.
                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                Faris M, Cha
Jakobovits A;
                                                                                                                                                                                                                                                                                                                     29-NOV-2001.
                                                                                                                                                                                                                                      AAS99443;
                                                                     products.
                                                                                                                                                                                                                    RESULT 372
AAS99443/c
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The invention relates to an isolated 98P7C3-related protein which is a homeodomain protein highly expressed in various cancers. Also include are homeodomain protein highly expressed in various cancers. Also include are polymucleotides encoding the protein or proteins 90% identical to 98P7C3, a pharmaceutical composition comprising the polymucleotides (including an expression vector comprising the 98P7C3 encoding polymucleotides) or a comparation or polymucleotides in a biological ample, monitoring the 98P7C3 protein or polymucleotides in a biological sample, monitoring the 98P7C3 protein or polymucleotides and a pharmaceutical composition comprising a modulator of 98P7C3. 98P7C3 protein, or T cell/B cell composition or polymucleotides and a pharmaceutical composition comprising a modulator of 98P7C3. 98P7C3. protein, or T cell/B cell composition in immunogens derived from it, are useful in inducing an immune response (in mammal) to a 98P7C3 protein. Upon contact with a cytotoxic T cell/B cell composition or epitope. The antibody is useful for delivering a cytotoxic agent to the antibody or its fragment that specifically binds cytotoxic agent to the antibody or its fragment that specifically binds conjugate. The modulator is useful for treating a patient with a cancer that expresses 98P7C3. by conjugation with a cancer, conjugate. The modulator is useful for treating a patient with a cancer conjugate. The modulator is useful for treating a patient with a cancer conjugate. The modulator such that the vector conjugate the modulator, such that the vector cells and exposing the method of suppression subtractive colls and the anced of an antibody coding sequence to the cancer cells and the encoded single chain antibody coding sequence to the cancer cells and the encoded single chain antibody coding sequence to the cancer cells and the encoded single chain antibody especies of that expresses of intracer and colon cancer), by administering the coll of an actual colon cancer of a cubh sequence for 98P7C3 by the method o
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                                                                                                                                                                                                      New isolated 98P7C3-related homeodomain protein highly expressed in various cancers, useful in cancer vaccines and for generating immune response directed to 98P7C3 in mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                    Faris M, Afar DEH, Levin E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                   Example 1; Page 53; 155pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 TCCTCGGCCGCGACCACG 3
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A;
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                                                    Challita-Eid PM, Hubert
Mitchell SC, Jakobovits
                                                                                                                                                 WPI; 2002-097642/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequences AAS95810-AAS95820 represent the 103P3E8 gene and the primers and adaptors used to amplify 103P3E8 DNA. 103P3E8 exhibits tissue and adaptors used to amplify 103P3E8 DNA. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 bladder, comprising a polymucleotide protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P3E8 related protein, and a ribozyme capable of cleaving a polymucleotide having the 103P3E8 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P3E8. The sequences can be used in diagnostic methods to monitor the level of 103P3E8 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                        Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, 98P6G3; ss; homeodomain protein; vaccine; cytostatic. epitope; transgenic animal; immunogen; T cell; B cell; cytotoxic T cell; CTL; prostate cancer; bladder cancer; kidney cancer; lung cancer; breast cancer; uterine cancer; cervical cancer; stomach cancer; rectal cancer; colon cancer; chromosome 4q11-q12; PCR primer; adapter; suppression subtractive hybridisation; SSH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human cancer related protein 98P7C3 nested PCR primer 2.
                                                                                                                                                                                                                Raitano AB, Mitchell SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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Gaps

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JP2001321190-A. Homo sapiens.

20-NOV-2001

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Claim 4; Page 18; 528pp; Japanese.
   12-MAR-2001; 2001JP-00068285.
           10-MAR-2000; 2000JP-00066716.
                    (RIKA ) RIKAGAKU KENKYUSHO (GENO-) GENOTEX YG.
                                        Arraying genome clones.
                                WPI; 2002-144136/19.
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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant correlating the clones having said marker sequence; (d) the multiwell copies containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. or array the multiwell copies; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell copies; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell completes; (e) the mixed clones are cultured and the call lateral directions; (f) the mixed clones are cultured and the creatitate cultures are amplified by using the above primer; (g) signals creatituted as the positions on the chromosome and arrayed. The clones are reconstituted as the positions on the chromosome and arrayed. The mixed plates are sepecified from the malysis. ABM45323 to ABM45324 represent propresent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;

Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels 357 AGCGACTICCICACTITC 374 1 AACGACTTCCTCACGGTC 18 g ò

0; Gaps

Human PTP1B mRNA level inhibition antisense DNA #1. ABK37204 standard; DNA; 20 BP. (first entry) 08-MAY-2002 ABK37204; RESULT 374 ABK37204 

Human; mouse; rat; protein tyrosine phosphatase 1B; PTP1B; ss; adipose; liver; kidney; metabolic disease; type 2 diabetes; obseity; cancer; hyperproliferative condition; blood serum; blood plasma; antidiabetic; blood glucose level; cytostatic; anorectic; antisense gene therapy; PTP1B mRNA level inhibition.

Homo sapiens

WO200210378-A2.

07-FEB-2002,

30-JUL-2001; 2001WO-US023874.

31-JUL-2000; 2000US-00629644.

(ISIS-) ISIS PHARM INC.

Novel antisense compound useful for treating type 2 diabetes, cancer and obssity, is targeted to nucleic acid encoding human protein phosphatase 18, and hybridizes and inhibits PTP18 expression. Cowsert LM, Wyatt J, Freier SM, Monia BP, Butler MM, Mckay R; WPI; 2002-180079/23.

Example 15; Page 67; 142pp; English.

The invention relates to a compound targeted to a nucleic acid molecule encoding procein phosphatase 1B (PTP1B), which specifically hybridises with and inhibite the expression of PTP1B. The compounds of the invention are useful for inhibiting the expression of PTP1B in liver, kidney or adipose cells or tissues and for treating an animal, preferably human, having a disease or conditions associated with PTP1B, including metabolic hyperproliferative conditions such as cancer. The sequences are also useful for decreasing blood (serum or plasma) glucose levels in an animal e.g. a diabetic human or rodent, for preventing or delaying the onset of a disease or condition associated with PTP1B, and for preventing or delaying the onset of an increase in blood glucose levels. This sequence represents a PTP1B mRNA level inhibition antisense oligonucleotide of the invention 

Sequence 20 BP; 2 A; 11 C; 5 G; 2 T; 0 U; 0 Other;

Gaps ô Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels

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303 CTGAGCCCCGGGGACCGC 320 

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ABK37412 standard; DNA; 20 BP. 08-MAY-2002 (first entry) ABK37412; **ABK374** 

RESULT 375

Human; mouse; rat; protein tyrosine phosphatase 1B; PTP1B; ss; adipose; liver; kidney; metabolic disease; type 2 diabetes; obesity; cancer; hyperproliferative condition; blood serum; blood plasma; antidiabetic; blood glucose level; cytostatic; anorectic; antisense gene therapy; pTP1B mRNA level inhibition. Rat PTP1B mRNA level inhibition antisense DNA #129.

30-JUL-2001; 2001WO-US023874. Rattus norvegicus. WO200210378-A2. 07-FEB-2002. 

31-JUL-2000; 2000US-00629644.

ISIS-) ISIS PHARM INC.

Mckay R; Wyatt J, Freier SM, Monia BP, Butler MM, WPI; 2002-180079/23. Cowsert LM,

Novel antisense compound useful for treating type 2 diabetes, cancer and obesity, is targeted to nucleic acid encoding human protein phosphatase 1B, and hybridizes and inhibits PTP1B expression.

8X866666666666668X8X

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73; 142pp; English
Claim 3; Page
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The invention relates to a compound targeted to a nucleic acid molecule encoding protein phosphatase 1B (PTP1B), which specifically hybridises with and inhibits the expression of PTP1B. The compounds of the invention are useful for inhibiting the expression of PTP1B in liver, kidney or adjoce cells or tissues and for treating a naimal, preferably human, having a disease or condition associated with PTP1B, including metabolic diseases or conditions, e.g. type 2 diabetes and obesity, or hyperproliferative conditions such as cancer. The sequences are also useful for decreasing blood (serum or plasma) glucose levels in an animal e.g. a diabetic human or rodent, for preventing or delaying the onset of an increase in blood glucose levels. This sequence represents a PTP1B mRNA level inhibition antisense oligonucleotide of the invention

Sequence 20 BP; 2 A; 8 C; 6 G; 4 T; 0 U; 0 Other;

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..
0
Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02;
                          3; Indels
                           0; Mismatches
                                                    270 CTGGAGCAGGGCGGCACC 287
                                                                          N
3.1%;
1 Similarity 83.3%;
15; Conservative (
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 Query Match
Best Local Similarity
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**ABK37400** RESULT

ABK37400 standard; DNA; 20 

BP.

(first entry) 08-MAY-2002 ABK37400;

Rat PTP1B mRNA level inhibition antisense DNA #117.

Human; mouse; rat; protein tyrosine phosphatase 1B; PTP1B; ss; adipose; liver; kidney; metabolic disease; type 2 diabetes; obesity; cancer; hyperproliferative condition; blood serum; blood plasma; antidiabetic; blood glucose level; cytostatic; anorectic; antisense gene therapy; PTP1B mRNA level inhibition.

Rattus norvegicus.

WO200210378-A2

07-FEB-2002

30-JUL-2001; 2001WO-US023874.

31-JUL-2000; 2000US-00629644.

(ISIS-) ISIS PHARM INC

Mckay Butler MM, Freier SM, Monia BP, Wyatt J, WPI; 2002-180079/23. Cowsert LM,

Novel antisense compound useful for treating type 2 diabetes, cancer and obesity, is targeted to nucleic acid encoding human protein phosphatase 1B, and hybridizes and inhibits PTPIB expression.

Example 16; Page 72; 142pp; English.

The invention relates to a compound targeted to a nucleic acid molecule encoding protein phosphatase 1B (PTP1B), which specifically hybridises with and inhibits the expression of PTP1B. The compounds of the invention are useful for inhibiting the expression of PTP1B in liver, kidney or adipose cells or tissues and for treating an animal, preferably human, having a disease or condition associated with PTP1B, including metabolic

diseases or conditions, e.g. type 2 diabetes and obesity, or hyperproliferative conditions such as cancer. The sequences are also useful for decreasing blood (serum or plasma) glucose levels in an animal e.g. a diabetic human or rodent, for preventing or delaying the onset of a disease or condition associated with PTHB, and for preventing or delaying the onset of an increase in blood glucose levels. This sequence represents a PTPIB mRNA level inhibition antisense oligonucleotide of the invention 8888888888

Sequence 20 BP; 2 A; 6 C; 8 G; 4 T; 0 U; 0 Other;

Gaps . 0 Length 20; 3; Indels Score 13.2; DB 1; Pred. No. 3.8e+02; 0; Mismatches 3.1%; Local Similarity 83.3%; Les 15; Conservative Query Match Best Loca Matches

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ВЪ. ABT12959 standard; DNA; 20

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Gaps

ABT12959;

(first entry) 17-JAN-2003 Mycobacterium tuberculosis-specific DNA sequence #46.

Mycobacterium detection method; PCR; primer; probe; ss.

Mycobacterium tuberculosis.

WO200274991-A2

26-SEP-2002.

20-MAR-2002; 2002WO-GB001308

20-MAR-2001; 2001GB-00006949

(NORC-) NORCHIP AS. (ALLA/) ALLARD S J.

Karlsen F;

WPI; 2002-750564/81.

Detecting the presence of Mycobacterium tuberculosis in a test sample, comprises inducing mRNA expression of Mycobacterium tuberculosis and detecting the induced mENA.

Claim 8; Page 14; 70pp; English.

The invention comprises a method for detecting the presence of a microorganism (particularly Mycobacterium tuberculosis) in a test sample. The method of the invention comprises exposing the test sample to an inducer that is capable of inducing the expression of at least one gene in the micro-organism and then testing for the presence of mRNA from this gene. The method of the invention is useful for detecting an mRNA that is tuberculosis). The present DNA sequence represents a Mycobacterium tuberculosis). The present DNA sequence represents a Mycobacterium specific nucleotide which can be used as a primer or probe in the method the invention 

Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ive 0; Mismatches 3; Indels 3.1 Best Local Similarity 83.3 Matches 15; Conservative

ö

Gaps

374 CCTGGACCGCGACGACGG 391

8

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Page 182
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Antisense PCR primer used to amplify GAPDH cDNA
                                                                                                                     JS2002107219-A1.
                                                                                               Homo sapiens
                                                                                                                                           08-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       요
  ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention comprises antisense oligonucleotides designed to inhibit expression of Syntaxin 4 interacting protein. The antisense oligonucleotides of the invention are useful for inhibiting the antisense oligonucleotides of the invention are useful for thibiting the antisense oligonucleotides are also useful for treating or tissues. The antisense or condition associated with Syntaxin 4 interacting protein (e.g. diabetes, obesity or a skeletal muscle disorder). The antisense oligonucleotides can also be used to prevent or delay infection, inflammation and tumour formation. The present DNA sequence represents a human Syntaxin 4 interacting protein antisense oligonucleotide. NoTE: The methoxyethyl wings
                                                                                                                                                                  Human, antisense gene therapy, Syntaxin 4 interacting protein, ss; antisense oligonucleotide; diabetes; obesity; skeletal muscle disorder; inflammation; tumour formation; phosphorothioate backbone; 2'-0-methoxyethyl wing.
                                                                                                                                                                                                                                                                                                                                                                                                                Novel antisense compound that hybridizes and inhibits nucleic acid molecule encoding Syntaxin 4 interacting protein, useful for treating diabetes, obesity and skeletal muscle disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                             Human syntaxin 4 interacting protein antisense oligonucleotide 54,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 4 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 84; 154pp; English.
                                                                                                                                                                                                                                                                                                                                                                      Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   116 CAGCAAGTACGGCATGCT 133
2 CCTGGACTGGGACTACGG 19
                                                                       BP.
                                                                                                                                                                                                                                                                                                  19-SEP-2001; 2001WO-US029251.
                                                                                                                                                                                                                                                                                                                         22-SEP-2000; 2000US-00668313
                                                                       ABQ62315 standard; DNA; 20
                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                              ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                      Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-404952/43.
                                                                                                                                                                                                                                                    WO200224864-A2.
                                                                                                                     16-AUG-2002
                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                          28-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                      Monia BP,
                                                                                              ABQ62315;
                                                 RESULT 378
                                                            ABQ6231
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The invention discloses an adenoviral vector for the selective expression of a toxin gene in a cancer cell. The toxin gene is operably linked to a promoduce of a gene with undetectable expression in liver, so that promoduce of a gene with undetectable expression in liver, so that expression of the toxin gene is reduced in liver cells. Adenoviral vectors and adenoviral gene therapy have been used to introduce suicide/toxic genes to advanced gastrointestinal or pencreatic cancer cells. This technique has a problem due to the hepatotropism of the adenovirus for systemically administered adenoviral vectors localise corringally to the liver, where the suicide gene therapy of intrahapatic tumour leads to severe liver dysfunction. Cyclooxygenses-2 (Cox-2) has virtually undetectable expression in most tissues, but is closely linked to carcinogenesis and progression of colon cancers. The promoter of Cox-2 therefore has a tumour "on" liver "off" expression profile, which can be utilised in the adenoviral gene therapy vectors. The adenovirus is used to illised in the adenovirus gene therapy particularly concerted to suplify glyceraldehyde-3-concerted to amplify gly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 An adenoviral vector containing a toxin gene under control of a promoter with undetectable expression in the liver is useful to treat gastrointestinal or pancreatic cancer with reduced liver toxicity.
PCR; primer; ss; adenoviral vector; toxic gene; cancer; promoter; liver; gene therapy; suicide gene; gastrointestinal cancer; pancreatic cancer; hepatctropism; adenovirus; interbepatic tumour; liver dysfunction; cyclooxygenase-2; Cox-2; carcinogenesis; colon cancer; tumour; liver toxicity; GAPDH; glyceraldehyde-3-phosphate dehydrogenase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ó
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            profile of cyclooxygenase-2 (Cox-2) cDNA in various tissues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human BH3 interacting domain death mRNA agonist inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 4 A; 7 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 4; 35pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        85 CAGTGGACATCACCACGT 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 CAGTGGACTCCACGACGT 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-DEC-2001; 2001US-00005964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-DEC-2000; 2000US-0251375P
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(first entry)
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Query Match
Best Local Similarity 83.3%
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Curiel DT, Yamamoto M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-697880/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURI/) CURIEL D T. (YAMA/) YAMAMOTO M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-2003
15-AUG-2002
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caacaagragracardcr

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ABX14127 standard; DNA; 20

RESULT 379

ABX14127

(first entry)

27-FEB-2003

ABX14127;

SAXAXEX

Hepatotrophic; immunomodulatory; cytostatic; antiinflammatory; hepatitis; haemostatic; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl; 2'-MOE; phosphorothioate backbone; ds.

Homo sapiens

Chimeric

WO200220547-A1.

14-MAR-2002

31-AUG-2001; 2001WO-US027316

07-SEP-2000; 2000US-00657346. 07-MAR-2001; 2001US-00800631.

(ISIS-) ISIS PHARM INC.

Wyatt JR; Zhang H, WPI; 2002-393838/42.

Novel antisense compound targeted to nucleic acid molecule encoding the BH3 interacting domain death agonist, useful for treating animals with diseases associated with BH3 interacting domain death agonist, e.g. hepatitis.

Claim 3; Page 87; 171pp; English.

The invention relates to a compound 8 to 50 nucleotides in length
targeted to a nucleic acid molecule encoding a BH3 interacting domain
death agonist, where the compound specifically hybridses with and
inhibits the expression of the BH3 interacting domain death agonist. The
compound of the invention is useful for inhibiting the expression of the
BH3 interacting domain death agonist in cells or tissues. The compound is
also useful for treating an animal having a disease or condition
associated with the BH3 interacting domain death agonist, e.g.
haematopoietic disorder, hyperproliferative disorder, a developmental
disorder, immunological disorder, or a disease or condition of the liver
c e.g., hepatitis, or a condition associated with apoptosis. The compound
is useful for diagnostics, therapeutics, proph/laxis and as research
c reagents and kits. This polynucleotide sequence represents an antisense
c oligonucleotide inhibitor of the DNA from human BH3 interacting domain
death agonist RNA of the invention. NOTE: This sequence is a chimeric
c oligonucleotide 20 nucleotides in length, which is flanked on both sides
by five-nucleotides. The wings are composed of 2'-methoxyethyl (2'
"NOE) nucleotides. The internucleoside (backbone) linkages are
c phosphorothoate (P=S) throughout the oligonucleotide. (Updated on 29-AUG 

Sequence 20 BP; 4 A; 4 C; 10 G; 2 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; live 0; Mismatches 3; Indels Best\_Local Similarity 83.3 Matches 15; Conservative Query Match

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114 CGCAGCAAGTACGGCATG 131 **8** CGGAGCAAGGACGGCGTG

Б ò

B.

Human C/EBP beta phosphorothioate antisense oligonucleotide, SEQ ID:40.

MOE MOE Human, C/BBP beta, CCAAT/enhancer-binding protein beta, C/BPB; LAP; TCF5, CRP2; NRIG; IL6DBP, NR-M, AGP/BBP; Apc/BBP; transcription factor; tissue development; cellular inction; proliferation; differentiation; hormone responsivenes; oxidative stress respons.

IL-6 signalling mediator; interleukin-6; carbohydrate metabolism; immunity; Thi response; female fertility; gluconeogenesis; ovarian; cancer; tumour formation; type II; diabetes; infection; inflammation; expression inhibition; phosphorothioate; antisense oligonucleotide; ss. 'n /note= "2'-methoxyethyl (2'-MOE) nucleotides. All 2' cytosines are 5-methylcytosine" /\*tag= c /\*tag= c /\*mad base= OTHER //note= "2'-methoxyethyl (2'-WOE) nucleotides. All cytosines are 5-methylcytosine" Location/Qualifiers /mod\_base= OTHER 1. .5 /\*tag= b 16. .20 Key modified\_base modified\_base modified\_base Homo sapiens.

US6271030-B1

07-AUG-2001.

14-JUN-2000; 2000US-00593711.

14-JUN-2000; 2000US-00593711.

(ISIS-) ISIS PHARM INC

Butler MM, Wyatt J; Monia BP,

WPI; 2002-214451/27.

Novel antisense compound targeted to nucleic acids encoding human or mouse CCAAT/enhancer binding protein (C/EBP) beta, useful in vitro for inhibiting expression of human or mouse C/EBP beta in cells/tissues.

Claim 1; Col 42; 69pp; English.

Sequences AB194252-AB194476 represent antisense oligonucleotides targeted to the human or mouse CCAAT/enhancer-binding protein alpha (C/BBP alpha) and which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human and/or mouse C/BBP alpha RNA, and were analysed for their effect on C/BBP alpha mRNA levels of granditation factors which regulate the expression of a wide range of transcription factors which regulate the expression of a wide range of conditions and functional differentiation. C/BBP beta (also Known as C/BBB2, LAP, TCF5, CRP2, NFILG, ILGOBP, NFW, AGP/BBP and Apc/BBP) conditions and functional differentiation. C/BBP beta (also Known as C/BBB2, LAP, TCF5, CRP2, NFILG, ILGOBP, NFW, AGP/BBP and Apc/BBP) conditions a mediator of IL-6 (interleukin-6) signalling. C/BBP beta is thought to be involved in carbobydate metabolism, immunity, the Th1 cresponse, female fertility and gluconeogenic pathways. C/BBP beta is compared in malignant ovarian tissue compared with normal ovarian tissue. Conditions and testais, with the conditions and itsue compared with normal ovarian tissue. Conditions associated with C/BBP beta as indicating that it is involved in the conditions are useful for diagnosis, prevention and treatment of impairment of insulin secretion in type II diabetes. The oligonucleotides conditions associated with C/BBP beta expression, guch as cancer conditions associated with C/BBP beta expression, guch as cancer conditions undertes), infection, or inflammation.

4 T; 0 U; 0 Other; ა ც A; 9 C; BP; 2 20 Sequence

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The invention relates to monitoring 83P2H3 (a calcium transport protein whose gene is located on chromosome 7q34) gene products in a biological eample from a parient who has or is suspected of having cancer cancer), computed (a) determining the status of 83P2H3 gene products expressed by calls in a tissue sample from an individual and (b) comparing the status to the status of 83P2H3 gene products in a normal sample. Also included are modulators of 83P2H3 function or status, generating antibodies/immune response against 83P2H3 conjugating the agent to an entigen) binding peptides derived from the protein delivering a cytotoxic agent to a cell cypridma that produces the recombinant protein, a non-human transgenic animal that produces the recombinant protein, a non-human transgenic animal that produces the recombinant protein, a non-human of the anti-83P2H3 antibody, a vector comprising a non-human of the anti-83P2H3 antibody, a vector comprising a non-human soft the anti-83P2H3 antibody, a vector comprising a non-human system T cell or B cell epitope, and contacting the protein that comprises a T cell or B cell respectively. The method is useful for monitoring the presence of cancer in an individual. The modulator is cancer that expresses 83P2H3. The immunological methods are useful for cancer that expresses 83P2H3. The immunological methods are useful for Monitoring 83P2H3 gene products for monitoring the presence of cancer in a subject, comprises determining the status of 83P2H3 gene products in a tissue sample from the subject and comparing it to a normal sample. Gaps Human; human leukocyte antigen; HLA; immunogen; 83P2H3; CaTrF2E11; calcium transport protein; cancer; prostate cancer; cytostatic; chromosome 7q34; chromosome 12q24.1; T cell; B cell; Bs; primer. ; 0 Afar DEH; Length 20; Query Match 3.1%; Score 13.2; DB 1; Length 2
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels Raitano AB, Challita-Eid PM, Faris M, Saffran DC, Levin E, Hubert RS, Ge W, Jakobovits A; Human 83P2H3 cDNA isolation nested PCR primer 2. Example 1; Page 76; 270pp; English. 146 GGTGGAGGCCGGCTTCGA 163 18 GGCGGAGCCGGGCTTCGA 1 BP 17-AUG-2001; 2001WO-US025782 17-AUG-2000; 2000US-0226329P 422/c ABK67422 standard; DNA; 20 02-JUL-2002 (first entry) WPI; 2002-269179/31. (AGEN-) AGENSYS INC. WO200214361-A2. Homo sapiens. 21-FEB-2002. ABK67422; RESULT 38 ABK67422/ 셤 ઠ

ö generating an immune response against 83P2H3, and for detecting the presence of 83P2H3-related protein or polynucleotide in a biological sample from a patient who has or who is suspected of having cancer. The antibody is useful in prostate cancer diagnosis, prognosis, imaging methodologies and treatment, to detect and quantify 83P2H3 imaging solating 83P2H3 homologues/related molecules, and for generating idiotypic antibodies that mimic the 83P2H3 related protein, for idiotypic antibodies that mimic the 83P2H3 protein. The present sequence is a PCR primer used in the isolation of cDNA encoding 83P2H3 or its related protein CaTrF2EH1 Gaps ö Human, cytostatic, 85P1B3; cancer; immunogen; ss; primer; PCR; chromosome 15q14. Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; Human cDNA 85P1B3 nested PCR primer 2. 373 TCCTGGACCGCGACGACG 390 TCCTCGGCCGCGACCACG 3 ABK70514 standard; DNA; 20 BP. 28-AUG-2001; 2001WO-US026838. 28-AUG-2000; 2000US-0228432P. 15-JUL-2002 (first entry) (AGEN-) AGENSYS INC. WO200218578-A2. Homo sapiens. 07-MAR-2002. ABK70514; 20 RESULT 383 888888888888 셤 ሯ

encoding, Composition for modulating the status of 85P1B3 protein or a molecule comprising a substance e.g. antibody specific to, nucleic acid encodir or ribozyme of 85P1B3. WPI; 2002-382963/41.

Ge W, Challita-Eid P;

Faris M, Hubert RS, Afar D,

Raitano AB, I Jakobovits A;

Example 1, Page 76; 201pp; English.

The invention relates to a composition comprising a substance that
modulate the status of 85PBB3, where the status of a cell expresses
85PBB3 gene product is modulated. Also included are a composition
comprising a peptide region of 5 amino acids of the 85PBB3 protein, in
any whole number increment up to 229 that includes an aa position
selected from an aa position having a value greater than 0.5 in the precent accessible residue profile, an aa position having a value greater than
0.5 in the percent accessible residue profile, an aa position having a value greater than
0.5 in the average flexibility profile; a an aa position having a value greater than
0.5 in the average flexibility profile; a position having a value greater than 0.5 in the average flexibility profile; a position having a value greater than 0.5 in the beta-turn profile; a polymucleotide that encodes analogue peptide of 8, 9, 10 or 11 contiguous residues of the 85PBB3 protein; a recombinant protein comprising the antigen-binding region of a monocloral antibody; a non-human transgenic animal that produces an antibody that binds to the 85PBB3 protein; a

hybridoma that produces antibody specific to the protein; a single chain monoclonal antibody specific to the variable domains of the heavy and monoclonal antibodies specific to the protein; a vector comprising a polymucleotide that econes the MAD; inhibiting growth of cancer cells or treating a patient who bears cancer cells that expresses the protein, antibody, polymucleotide cancer cells that expresses ribozyme that cleaves the polymucleotide to the polymucleotide or ribozyme that cleaves the polymucleotide and T cells that specifically risozyme that cleaves the polymucleotide and T cells that specifically directed to the protein and generating a mammalian immune system to an immunogenic portion of the protein or polymucleotide. The composition, which comprises an antibody specific to the protein, is useful for a delivering a cytotoxic agent to a cell that expresses the protein by providing a cytotoxic agent to a cell that expresses the protein by providing a cytotoxic agent to call that expresses the protein by providing a cytotoxic agent to manipody and exposing the cell compared to the antibody agent to a cell that expresses the protein by providing a cytotoxic agent to acell that expresses the protein of compared to antibody and exposing the compared of the protein of contract cells or treating a mammalian immune response the protein, for generating a mammalian immune response of the protein, for detecting the presence of the protein or polymucleotide in a biological ample in a patient who has or who is suspected of having cancer and for monitoring 85P183 in a biological end on human chromosome 15914. The present

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ö Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels 373 TCCTGGACCGCGACG 390 20 recredecececeáceáce 3 셤 ઠે

RESULT ABI9363

ABI93630 standard; DNA; 20 BP CXSXTXEXEXEXEXEXEXEXEXEXEXEXEXEXEXEXC

(first entry) 15-FEB-2002 Capture oligonucleptide Zip ID#717 oligo #9.

Human, K-ras, PCR primer, probe, capture probe, mutation detection, ligase detection reaction, LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forenaic; environmental monitoring; food industry; feed industry; ss.

Synthetic

WO200179548-A2.

25-OCT-2001

14-APR-2000; 2000US-0197271P.

04-APR-2001; 2001WO-US010958.

CORR ) CORNELL RES FOUND INC.

Favis R, Gerry NP, Zirvi M, Barany F,

Kliman R;

WPI; 2002-034366/04.

Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.

Example 5; Fig 29; 300pp; English.

The present invention describes a method (M1) for designing capture

cc oligonucleotide probes (I) for use on a support to which complementary cligonucleotide probes (II) will hybridise with little mismatch, where cligonucleotide probes (II) will hybridise with little mismatch, where cc (I) have melting temperatures within a narrow trange. The method is useful cfor detecting infectious diseases caused by bacterial infectious agents cc s.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents corrected fungatutus, viruses e.g. T-cell lymphocytotrophis citus, Epstein-Barr virus and pollo virus, and parasitic infectious agents cc selected from Onchoverva volvulus, Entamoeba histolytica and Dracumculus cas 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

Companies of the method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

Concerting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, prover oused for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the companies of particular sites and indentifying if ligation of the oligonucleotide probe companies or absence of the target nucleotide sequences. ABI82074 to represent oligonucleotide sequences used in the exemplification of the present on a particular sites. the present invention \$

Sequence 20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;

. 0 Length 20; 3; Indels Score 13.2; DB 1; Pred, No. 3.8e+02; 0; Mismatches 3; 3.1%; Best Local Similarity 83.3%; Matches 15; Conservative (

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29 GGGCTGGGACGAAGATGG 46

ò d

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Gaps

RESULT 385 AB1929

ABI92926 standard; DNA; 20

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AB192926;

15-FEB-2002 (first entry)

Capture oligonucleptide Zip ID#13 oligo #9.

Human, K-ras, PCR primer, probe, capture probe, mutation detection, ligase detection reaction, LDR, p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome, obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forenaic; environmental monitoring; food industry; feed industry; ss. 

Synthetic.

WO200179548-A2.

25-OCT-2001.

04-APR-2001; 2001WO-US010958.

14-APR-2000; 2000US-0197271P.

FOUND INC. (CORR ) CORNELL RES Gerry NP, Favis R, Barany F, Zirvi M,

Kliman R;

WPI; 2002-034366/04

to which Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch

Example 5; Fig 29; 300pp; English.

The present invention describes a method (M1) for designing capture

Wed Apr 21 12:58:21 2004

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cc oligonucleotide probes (I) for use on a support to which complementary cligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful cc. salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Canida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polito virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolyrica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obseity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, cancer is specifically associated with a gene selected from BRCA1 gene, psi gene, human papillomavirus types 16 and 18 and liver cancers. The cancer is specifically detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the computer, detecting dentifying if ligation of the oligonucleotide probe sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences ABI82074 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of the present invention
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Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 4 C; 8 G; 5 T; 0 U; 0 Other; 112 ACCGCAGCAAGTACGGCA 129

19 ATCGCTGCAAGTACCGCA 2 a

AAL40496 standard; DNA; 20 BP. RESULT 386 AAL40496/c

AAL40496;

Cytostatic, 158P1D7; cancer, bladder cancer, mouse, rat; rabbit; dog; cat; cow; horse; human; vaccine; gene therapy; PCR; primer; ss. 158P1D7 cDNA related PCR primer SEQ ID No 668. 19-SEP-2002 (first entry) 

Homo sapiens.

22-AUG-2001; 2001WO-US026276. 22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P. WO200216593-A2. 28-FEB-2002.

(AGEN-) AGENSYS INC.

Levin E;

WPI; 2002-425659/45.

Faris M, Hubert RS, Raitano AB, Afar DBH, Challita-Eid PM, Jakobovits A;

New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horses or numans)

Example 1; Page 68; 181pp; English.

The invention relates to a novel nucleic acid, designated 158P1D7. The compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mice, rats, rabbits, dogs, cats, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This polynucleotide sequence represents a PCR primer of the 158P1D7 CDNA of the invention 88888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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AAL53476 standard; DNA; 20 BP. RESULT : 

Zinc transporter protein 108P5H8 nested primer 2.

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Gaps

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16-JAN-2003 (first entry)

Cytostatic; gene therapy; vaccine; zinc transporter protein 108P5H8; cancer; breast; colon; ovarian; lung; humoral; cellular immune response; passive immunisation; PCR; primer; ss.

Unidentified

WO200260953-A2. 08-AUG-2002.

17-DEC-2001; 2001WO-US049133.

15-DEC-2000; 2000US-0256210P. (AGEN-) AGENSYS INC.

Hubert RS, N Jakobovits A; Challita-Bid PM, Faris M, Afar DEH, Levin E, Morrison KJM, Raitano AB,

Mitchell SC;

WPI; 2002-627469/67.

Composition comprising a substance that modulates the status of a zinc transporter protein (108P5H8), useful in diagnosing and treating patients with cancer that express 108P5H8, such as breast, colon, ovarian or lung cancer.

Example 1; Page 95; 309pp; English.

The invention relates to a new composition comprising a substance that modulates the status of a zinc transporter protein, designated as modulates. The composition is 108P5H8, or a molecula that is modulated by 108P5H8. The composition is useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 108P5H8, such as breast, colon, ovarian or lung cancer. The 108P5H8 ene or its fragment can be used to elicit a humoral or cellular immune response. The antibodies are useful in active or passive immunisation. The 108P5H8 polymucleotides are useful in active and primers for the amplification or detection of 108P5H8 genes, as coding sequences for directing the expression of 108P5H8 genes. The polymucleotides of the invention can be used to treat disorders by gene therapy. This polymucleotide sequence represents a zinc transporter protein 108P5H8 related PCR primer of the invention

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83.3%;

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Afar DEH, Saffran D, Morrison K, Morrison RK, Ge W, Jakobovits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; 121P2A3; cytostatic; immunostimulant; vaccine; PCR; primer; humoral immune response; ss; callular immune response; ss; suppression subtractive hybridisation; SSH.
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                                                                       Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
0; Mismatches 3; Indels
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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human 121P2A3 post-SSH nested PCR primer 2.
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                                           3.1%; Scor.
83.3%; Pred
                                                                                                                                                                                                                              373 TCCTGGACCGCGACGACG 390
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25-APR-2001; 2001US-0286630P.
22-JUN-2001; 2001US-0300373P.
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                                                           Query Match
Best Local Similarity 83.3
Marches 15; Conservative
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DB 1; Length 20;

3.1%; Score 13.2;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a new composition comprising a substance that modulates the status of 158P3D2 or a molecule that is modulated by 158P3D2, where the status of a cell that expresses 158P3D2 is modulated. The composition is useful for treating cancer. This sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New composition comprising a substance that modulates the status of 158P3D2 or a molecule that is modulated by 158P3D2, useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, antisense, lung dysfunction; nasal airway dysfunction;
antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
            Gaps
                                                                                                                                                                                                                                      158P3D2; cytostatic; gene therapy; vaccine; cancer; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Hubert RS;
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            Indels
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Afar DEH, Ge W, Raitano AB, Challita-Eid PM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
 Pred. No. 3.8e+02;
                                                                                                                                                                                                               Novel protein 158P3D2 associated primer #4.
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                                      373 TCCTGGACCGCGACG 390
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25-APR-2001; 2001US-0286630P.
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Best Local Similarity 83.3
Matches 15; Conservative
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               15; Conservative
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                          RESULT 389
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antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, 1 Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 13919; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intitation codon, coding region, 5. or 3. end genomic flanking regions, 5. and 3. intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or neasl alraway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, companyepressive, and cytostaric activity. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a core preventing a respiratory lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing or depleting levels of, uniquinone or lung surfactant in a subject of the cathoring bronchocilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence at for sequences or condition.

Condition the printed prophylactic fersion or represented in the printed specification, but was obtained in electronic format directly from WIPO contents.

Sequence 20 BP; 3 A; 7 C; 9 G; 1 T; 0 U; 0 Other;

ö Gaps ö Cuery Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels

308 CCCCGGGGACCGCGTGCT 325 18 CCCCGGGGATGCCGTGCT

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ABZ86355 standard; DNA; 20 ABZ8635 ID ABZ8635 XX ABZ8635 XX ABZ8635 XX ABZ8635 XX YX Human C XX Human C

ABZ86355;

35

(first entry) 17-0CT-2003 Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic;

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Ka , Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

claim 15; SEQ ID NO 1597; 872pp; English.

first active agent comprising an oligonucleotide antisense to the invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubjudinone. A composition of the invention has antinflammatory, antiallergic, antisthatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a creventing a respiratory, lung or malignant disease or condition, also perventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing levels of adenosine of of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchocdilation, increasing levels of adenosine receptor, unug allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fire, wipo.int/pub/published\_pot\_sequences 

Sequence 20 BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

Gabs ö / Match 3.1%; Score 13.2; DB 1; Length 20; Local Similarity 83.3%; Pred. No. 3.8e+02; nes 15; Conservative 0; Mismatches 3; Indels Query Match Best Loca Matches

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ABZ99369/c ID ABZ99369 standard; DNA; 20 BP. ABZ99369;

RESULT 392

17-OCT-2003 (first entry) ZZXZXZXZX Z

Human PDE4C oligonucleotide sequence.

Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;

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antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. 

Homo sapiens

WO200285308-A2

31-OCT-2002

23-APR-2002, 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar Pabalan J, Li Y, Sandrasagra A, Katz E, Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure, SEQ ID NO 14611, 872pp, English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intitation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an intilnflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antisethmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entishery steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchocanstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed appearance of the control of the printed and the control of the sequence date for this patent is not represented in the printed and the control of the control of the sequence date for this patent is not represented in the printed and the control of the control of the sequence date for this patent is not represented in the printed and the control of the contr at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 1 A; 8 C; 8 G; 3 T; 0 U; 0 Other;

Gaps ; 0 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; Live 0; Mismatches 3; Indels 15; Conservative Local Similarity Query Match Best Loca Matches

297 AAGGACCTGAGCCCCGGG 314 ო

AGGACCTGAGCCCGCGG 20

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RESULT 393 ABZ92374
ID ABZS
XX
AC ABZS
XX
DT 17-C
XX
XX
DE Hume
XX
XX
KW Hume
KW anti

BP ABZ92374 standard; DNA; 20

ABZ92374;

(first entry)

17-0CT-2003

Human oligonucleotide sequence,

Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory, antiallergic;

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-0CT-2002

23-APR-2002, 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

Katz E, Pabalan J, S; Li Y, Sandrasagra A, Tang L, Shahabuddin (EPIG-) EPIGENESIS PHARM INC Nyce JW, I Miller S,

Aguilar

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure, SEQ ID NO 7616, 872pp, English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligomucleotide antisense to the initiation codon, coding region, 5' or 3' or and genomic flanking regions; 5' and 3' intron-exon junctions, 5' or 3' or region within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antistense gene therapy. The composition is useful for treating or preventing a respiratory, lung or mallignant disease or condition, as 1so for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodialation, increasing levels of adenosine receptor, producing bronchodialation, increasing levels of adenosine receptor, producing bronchodialation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease; or condition or lung inflammation, lung abeten is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences 

Sequence 20 BP; 5 A; 5 C; 7 G; 3 T; 0 U; 0 Other;

. 0 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; tive 0; Mismatches 3; Indels Local Similarity 83.3 nes 15; Conservative Query Match Best Loca Matches

27 GAGGGCTGGGACGAGAT 44

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RESULT 394 ABZ97852 ID ABZ9785

BP ABZ97852 standard; DNA; 20

ABZ97852;

(first entry) 17-0CT-2003

Human eotaxin oligonucleotide sequence.

Human, antisense, lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; \*\*\*\*\*

ubiquinone. 

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or gene therapy; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy, respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Katz E, Pabalan J, Aguilar Li Y, Sandrasagra A, K Tang L, Shahabuddin S; 24-APR-2001; 2001US-0286137P. 23-APR-2002; 2002WO-US013135. EPIG-) EPIGENESIS PHARM INC WPI; 2003-229219/22. WO200285308-A2. Homo sapiens. 31-OCT-2002, Nyce JW, I Miller S,

Disclosure; SEQ ID NO 13094; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5, or 3 end genomic flanking regions, 5, and 3 intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal alwayd yfsfunction and a second active agent comprising an antiinflammatory steroid and ubjunione. A composition of the invention has antiinflammatory, antiallargic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of ubjunione or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung alleraties, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in a leactronic format directly from WIPO or at fib., wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 1 A; 8 C; 4 G; 7 T; 0 U; 0 Other;

Gaps ö Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

308 CCCCGGGGACCGCGTGCT 325 CCCCTGGGACCTCGTTCT 19

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ABZ86905 standard; DNA; 20 ABZ86905; RESULT 399 ABZ86905/C XX ABZ80 XX DT 17-0 XX DE HUMB XX HUMB KW HUMB KW AURA

BP.

17-OCT-2003

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic;

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D; Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 2147; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intitation coodon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of imatiation codon, coding region, 5' or regions within 2-10 nucleotides of imatiation and a second active agent comprising an entiting all airway dysfunction and a second active agent comprising an entiting lammatory steroid and ubjquinone. A composition of the invention has antiinflammatory antiallargic antiasthmatic, hypotensive, also immunosuppressive, and cytostatic crivity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine or intifflammatory steroid in a subject, for reducing levels of adenosine ceptor, producing broncholation, increasing levels of unquinone or lung surfactant in a subject stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published\_pot\_esquences 

Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; llarity 83.3%; Pred. No. 3.8e+02; Conservative 0; Mismatches 3; Indels Local Similarity nes 15; Conserv Query Match Best Local Matches

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ACC47656 standard; DNA; 20 BP. ACC47656; RESULT 396 ACC47656

16-SEP-2003 (first entry)

Human IGFBPS phosphorothioate antisense oligonucleotide, SEQ ID NO:32.

Human; insulin-like growth factor binding protein 5; IGFBP5; IBP5; chromosome 2q33-34; IGF signal transduction; IGF regulation; apoptosis; \*\*\*\*

bone growth stimulator; hyperproliferative disorder; cancer; tumour; breast; prostate; pancreas; neuroendocrine; inflammatory disorder; colitis; developmental disorder; growth disorder; Duchenne muscular dystrophy; metabolic disorder; diabetes; osteoporosis; osteoperrosis; cytoetatic; antiinflammatory; expression inhibition; phosphorothicate; antisense oligonuclectide; ss. Location/Qualifiers Homo sapiens

1. .20 /\*tag= a Key modified\_base

/mod\_base= OTHER /note= "This oligonucleotide has a phosphorothicate /note= "This oligonucleotide has a phosphorothicate backbone and 2-'methyoxyethyl (2'-MOE) wings at the 5' and 3' ends, which are 5 nucleotides in length. Also all cytosine residues are 5-methylcytosines"

WO2003030826-A2

17-APR-2003

07-OCT-2002; 2002WO-US032060

09-OCT-2001; 2001US-00975123.

(ISIS-) ISIS PHARM INC.

Freier SM;

WPI; 2003-381673/36.

New antisense oligonucleotides for modulating insulin-like growth factor binding protein 5 gene expression, useful for preventing or treating cancers, inflammatory disorders, developmental disorders or metabolic disorders.

Claim 3; Page 76; 105pp; English.

Sequences ACC47637.ACC47667 represent phosphorothicate antisense cligonuclectides targeted to the human insulin-like growth factor binding protein 5 (IGFBPS) gane, which inhibit its expression. The antisense cligonuclectides were designed to target different regions of human coligonuclectides were designed to target different regions of human IGFBPS RNA, and were analysed for their effect on IGFBPS expression by quantitative real-time PCR. IGFBPS (also known as IBPS) is a member of the insulin-like growth factor superfamily, which are involved in the regulation of IGF action and bioavailablity, and which also mediate IGFP. Independent actions, including inhibition or enhancement of apoptosis. IGFBPS is a key component of the IGF system in bone, having a high specific binding affinity for hydroxyapatite and extracellular matrix of proteins, and appears to act as a growth factor, stimulating bone of proteins, and appears to act as a growth factor, stimulating bone of proteins, and appears to act as a growth factor, stimulating bone of proteins, and appears, notochord, muscle cells in a experimental model collis. It is also thought to play a role in prostate cancer of colitis. It is also thought to play a role in prostate cancer of colitis. It is expressed which high frequency in neuroendocrine tumours, and has been shown to be induced in breast cancer cells upon treatment with anticestrogens used to abolish tamoxifen resistant proliferation. The oligonucleotides of the invention and treatment of invention and treatment of information and treatment of information are useful for diagnosis, or prostate, pancreas of the invention and treatment of information and treatment of information and treatment of information and treatment of invention and treatment of invention and treatment of information and treatment of information and treatment of invention and treatment of information and treatment of invention and

Seguence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;

Gaps . 0 Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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CGAGGTCCTCAGTTTCCT 19 359 CGACTICCICACTITCCI

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ACA55326 standard; DNA; 20 BP ACA55326,

ACA55326;

05-JUN-2003 (first entry)

Human modified IgE CH2 domain PCR primer P171-A393

CH3 domain; IgE; antigen; non-anaphylactic; anti-IgE; fusion protein; dermatological; antiinflammatory; ophthalmological; allergy; asthma; allergy chinitis; gastrointestinal allergy; food allergy; eosinophilia; conjunctivitis; glomerular nephritis; flea allergy; atopic dermatitis; gene therapy; PCR; primer; ss.

Homo sapiens. Synthetic.

EP1262491-A2.

04-DEC-2002.

22-MAY-2002; 2002EP-00253606.

22-MAY-2001; 2001US-0292638P.

(PFIZ ) PFIZER PROD INC.

Brown IM, Morsey MA;

WPI; 2003-122561/12.

Novel isolated antigenic peptide comprising amino acid residues of CH3 domain of IgE molecule from first species and a second unrelated species, induces non-anaphylactic anti-IgE immune response in animal.

Example 2; Page 22; 50pp; English.

This invention describes a novel antigenic peptide comprising amino acid residues of an IgB CH3 domain from a first species (ADE1) and amino acid residues of an IgB CH3 domain from a first species (ADE2), where ADE1 is conserved in the IgE CH3 domain of the second apecies and ADE2 is not conserved in the IgE CH3 domain of the second species and ADE2 is not conserved in the IgE CH3 domain of the first species. The novel antigenic peptide induces a non-anaphylactic anti-IgE immune response in a nanimal. The invention also discloses the polynucleotide sequence conciding the antigenic peptide and an antigenic fusion protein comprising the antigenic peptide of the invention and a heterologous protein comprising the antigenic peptide of the invention and an enti-IgE immune response that does not cause anaphylaxis when administered to an animal. The carrier, where the fusion protein induces an anti-IgE immune response that does not cause anaphylaxis when administered to an animal. The products of the invention are useful in the manufacture of preventing IgE from binding to high affinity receptors on mast cells and operation are useful in the manufacture of a medicament for treating or preventing IgE-mediated allergic disorders including asthma, allergic rhinities, gastrointestinal allergic disorders consorted announced including asthma, allergic rhinities, gastrointestinal allergic disorders including asthma, allergic rhinities, gastrointestinal allergic disorders consorted products are useful for treating IgE-mediated allergic disorders by gene therapy. Antigenic peptides comprising conserved amino conditived by variable amino acid residues of the CH3 domain of an IgE molecule from as second unrelated species are capable of inducing a high titre of antitre of antibodies when administered to an animal without causing anaphylaxis. ACASIBH4-ACASIBH6 represent PCR primers used to amplify the polymucleotide sequences used in designing the constructs 

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The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding dual specific phosphatase 5, where the compound specifically hybridises with and inhibits the expression of dual specific phosphatase 5. The compound is used for treating an animal having a disease or condition associated with dual specific phosphatase 5 such as a hyperproliferative disorder, a developmental disorder, an inflammatory disorder or a disease which arises from aberrant apoptosis. Sequences ASK09162-ABX09139 represent human dual specific phosphatase 5 phosphorothioate oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; immunostimulant; 162P1E6; cytotoxic agent; immune response; cancer; bladder; prostate; kidney; lung; breast; passive immunisation; transgenic animal; vaccine; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense modulation of dual specific phosphatase 5 expression used in treating disorders e.g. inflammatory diseases.
                                                                                                                                                                                 Human; dual specific phosphatase 5; ss; developmental disorder;
hyperproliferative disorder; inflammatory disorder aberrant apoptosis;
antiinflammatory; cytostatic; antiapoptotic; antiproliferative;
phosphorothioate oligonucleotide.
                                                                                                                                              Human dual specific phosphatase 5 phosphorothioate oligonucleotide
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         063/c
ABX09063 standard; DNA; 20 BP.
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Best Local Similarity 83.3'
Matches 15; Conservative
                                                                                                           22-JAN-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-041418/03.
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                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                   ABX09063;
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XSXEXEXEXEXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New 151P3D4 proteins and genes, useful for eliciting a humoral or cellular immune response, or for diagnosing, prognosing, preventing or treating cancer, e.g. adenocarcinoma, bladder cancer, lung, breast cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; gene therapy; vaccine; modulator; 151P3D4; humoural; cancer; cellular immune response; adenocarcinoma; bladder; colorectal; lung; bronchial; breast; carcinoma; PCR; primer; ss.
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                                              Query Match
3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
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       Sequence 20 BP; 2 A; 6 C; 4 G; 8 T; 0 U; 0 Other;
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Morrison RK, Ge W, Jakobovits A;
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                                                                                                                                         384 GACGACGCCCCAAGAAG 401
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ABT43860 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                             DPNCDN nested primer 2 (NP2)
                                                                                                                                                                            GAGGACACCAAGAAG
                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2003 (first entry)
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Best Local Similarity
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WO200283860-A2

24-OCT-2002

Unidentified.

ABT43860;

ABT43860/

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RESULT 399

Best Loca Matches

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Gaps

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The invention relates to a novel composition comprising a substance that modulates the status of a 162P1E6 protein. The protein comprises one of 21 sequences of 70 - 146 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of a cell that expresses the protein is modulated. An antibody to the 162P1E6 protein is used to deliver a cytotoxic agent or a diagnostic agent to a cell that expresses the 162P1E6 protein. The composition is used to inhibit the growth of cancer cells or generate an immune response. The composition is used for detecting the presence of a 162P1E6 related protein or a learned or detecting them are useful for diagnosing, proteins and polymuclectide in a sample. The 162P1E6 proteins and polymuclectide in a bander cancer, prostate cancer, kidney cancer, lung cancer, such as bladder cancer. They can be used for eliciting a humoral or cellular immune response. The also be used for eliciting a humoral or cellular immune response. The passive immunisation. Transgenic animals are useful for developing and screening of useful reagents. The polymuclectide and polypetide screening of the invention can also be used to treat disorders by being used in a vaccine or in gene therapy. This polymuclectide sequence
                                                                                                                                                                                                                                                                                                                                                   Composition for diagnosing, prognosing, preventing or treating cancer, for eliciting a humoral or cellular immune response, or for active or passive immunization, comprises a substance that modulates the status of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STEAP-1; six transmembrane epithelial antigen of the prostate; cancer; cancer vaccine; delineation; cytogenetic abnormality; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Suppressive subtractive hybridisation of STEAP related primer #8.
                                                                                                                                                                                                                                           Challita-Bid PM, Raitano AB, Faris M, Hubert RS, Morrison K;
Morrison RK, Ge W, Jakobovits A;
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3.1%; Score 13.2; DB 1; Length 20;
Best Local Similarity 83.3%; Pred. No. 3.8e+02;
Matches 15; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 71; 437pp; English.
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ACD02621 standard; DNA; 20 BP.
                                                                                    09-APR-2002; 2002WO-US011544.
                                                                                                                               10-APR-2001; 2001US-0283112P
25-APR-2001; 2001US-0286630P
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                                                                                                                                                                                                                                                                                                             WPI; 2003-148268/14.
                                                                                                                                                                                                   (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                       a 162P1E6 protein.
WO200283916-A2
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                                            24-OCT-2002
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WO2003022995-A2.

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The invention describes a composition comprising a substance that modulates the status of a protein (1) of 340 or 283 amino acids, or of any of the 15 sequences of 259 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of the cell that expresses the protein is modulated. The compositions, proteins, polynucleotides and methods are useful for treating and detecting cancer. The STEAP-1-related proteins are useful for generating cancer vaccines. The polynucleotides are useful as tools for delineating, with greater precision, cytogenetic abnormalities in the chromosomal region that encodes STEAP-1 that may contribute to the malignant phenotype. This sequence represents a primer used to analyse human six transmembrane epithelial antigen of the prostate or STEAP-1 cDNA's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                               New composition comprising a substance that modulates the status of STEAP-1-related protein, useful for treating and detecting cancer.
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                                                                                                                                       Jakobovits A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; vaccine; cancer; immune response; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
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                                                                                                                                       Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jakobovits A, Challita-Eid PM, Faris M,
Morrison K, Morrison RK, Raitano AB;
                                                                                                                                                                                                                                               Example 1; Page 70; 248pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           °,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           373 TCCTGGACCGCGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ78176/c
ID ABZ78176 standard; DNA; 20 BP.
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10-APR-2001; 2001US-0283112P.
25-APR-2001; 2001US-0286630P.
                                                                                                                                       Raitano AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-APR-2002; 2002WO-US011654.
                             06-SEP-2002; 2002WO-US028371.
                                                         06-SEP-2001; 2001US-0317840P.
05-APR-2002; 2002US-0370387P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAY-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 83.3
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AGEN-) AGENSYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-075555/07.
                                                                                                                                                                     WPI; 2003-313240/30.
                                                                                                         (AGEN-) AGENSYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nested primer #2.
                                                                                                                                       Ge ₩,
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20-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ78176;
                                                                                                                                       Faris M,
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New composition comprising a substance that modulates the structure of proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune response in
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Example 1; Page 72; 1021pp; English

The present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful as probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the disposais and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, and as therapeutic agents. The proteins and peptides are useful as therapeutic, prognostic and diagnostic reagents for cancer. The present sequence is a primer, used in an example from the invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps . 0 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.88+02; 0; Mismatches TCCTGGACCGCGACGACG 390 Local Similarity 83.3 les 15; Conservative 373 Query Match Matches ઠ

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20 TCCTCGGCGCGACCACG 3 d

Cancer associated coding sequence PCR primer #3.

Cancer associated coding sequence; cancer; human; cytostatic; gene therapy; PCR; primer; ss.

10-APR-2002; 2002WO-US011645.

10-APR-2001; 2001US-0282739P. 10-APR-2001; 2001US-0283112P. 25-APR-2001; 2001US-0286630P. 10-APR-2002; 2002US-00286630P.

Challita-Eid

ö New pharmaceutical composition for diagnosing, prognosing, preventing of treating cancer, comprises a substance that modulates a nucleic acid sequence, e.g. 105P1B7, 152P1A2B or 156P3A6, or a molecule modulated by the nucleic acid.

The invention comprises the amino acid and coding sequence of a 184PIE2 protein. The DNA and protein sequences of the invention are useful for diagnosing, prognosing and/or treating cancer. The 184PIE2 DNA and protein sequences may also be used to elicit a humoral or a cellular immune response in patients and in monitoring genetic abnormalities. Antibodies raised against the 184PIE2 proteins may be use in active or passive immunisation. The present DNA sequence is used in the exemplification of the invention

Example 1; Page 69; 394pp; English.

Example 1; Page 34; 72pp; English.

The present invention relates to a pharmaceutical composition comprising a substance that modulates the status of a cancer associated mucheic acid sequence such as given in the specification (see ABZ20564-ABZ20575) or a molecule that is modulated by the above nucleic acid sequence, where the 

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Gape

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Query Match 3.1%; Score 13.2; DB 1; Length Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ö status of a cell that expresses the nucleic acid sequence is modulated. The composition is useful in diagnosing, prognosing, preventing and/or treating cancer. The nucleic acid sequence may be used in monitoring genetic abnormalities, in generating and characterising domain-specific antibodies, for identifying agents or cellular factors that bind to a protein, and in therapeutic and diagnostic contexts, such as diagnostic sasays, cancer vaccines, and methods of preparing vaccines. The present sequence is a primer used to identify the cancer associated coding sequences suitable to be modulated in the method of the invention New 184PIE2 polynucleotide encoding a 184PIE2 protein, useful for diagnosing, prognosing, preventing or treating cancer, in eliciting an immune response, and in chromosome mapping. Faris M, Hubert RS, Morrison K; ö Gene therapy, vaccine, 184P1E2, cancer, genetic abnormality; cellular immune response, immunisation, PCR, primer, ss. Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 184P1E2 gene-specific nested PCR primer #2. Chalitta-Eid PM, Raitano AB, Fari Morrison RK, Ge W, Jakobovits A; 373 TCCTGGACCGCGACGACG 390 20 TCCTCGGCCGCGACG 3 BP 10-APR-2001; 2001US-0282739P. 25-APR-2001; 2001US-0286630P. 09-APR-2002; 2002WO-US011643 Query Match 3.1%; Best Local Similarity 83.3%; Matches 15; Conservative AALS2254 Btandard; DNA; 20 (first entry) WPI; 2003-148269/14. (AGEN-) AGENSYS INC W0200283919-A2 Unidentified 16-0CT-2003 24-OCT-2002. AAL52254; RESULT 40 AAL52254/ 886666666666888 셤 ઠ

Length 20;

Query Match

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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding fibroblast growth factor (FGP) receptor 3 (also known as FGFR-3) ACH, UTK4 and CEK2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with FGFR-3 such as developmental disorders or hyperproliferative disorders, especially cancer of colorectal, bladder, bone, lung, cervical, breast or skin. They are useful as research reagents, therapeutics, prophylaxis, kits and diagnostics, and as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion of the genes expressed within cells and tissues. They are also useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human FGFR-3
                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Phosphorothioate backbone, All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                 Human, antisense; fibroblast growth factor receptor 3; prophylaxis;
developmental disorder; hyperproliferative disorder; antisense therapy;
FGFR-3; ACH; JTK4; CEK2; cancer; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel compound targeted to a nucleic acid molecule encoding fibroblast growth factor receptor 3, useful for inhibiting the expression of the receptor and for treating an animal having cancer or developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= b
/mod= base= OTHER
inod= "2'-methoxyethyl (2'-MOE) nucleotides"
in-.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod_base= OTHER
/note= "2 -methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                      Human FGFR-3 antisense oligonucleotide, ISIS #125169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 15; Page 79; 120pp; English
                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
/mod_base= OTHER
"Thosphor
373 TCCTGGACCGCGACGACG 390
                  465/c
AAD55465 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-SEP-2002; 2002WO-US028549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-SEP-2001; 2001US-00953047
                                                                                                                                                                       07-AUG-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-313244/30.
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modified_base
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                                                                                                                                                                                                                                                                                                                   Synthetic
                              20
                                                                                                                                        AAD55465;
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                                                                         RESULT 40
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Seguence 20 BP; 5 A; 6 C; 7 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding BLCL2-associated X (BAX) protein, where the compound specifically hybridises with the nucleic acid molecule encoding BAX protein and inhibits the expression of BAX protein. The compound specifically hybridises with at least 8-nucleobase portion of an active site on a nucleic acid molecule encoding BAX protein. Also described: (1) a composition of 13 and a pharmaceutical carrier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compounds, useful for modulating the expression of BCL2-sasociated X (BXX) protein or for treating a disease or condition associated with BAX protein, e.g. Parkinson's disease, Hodgkin's disease or Alzheimer's disease.
                                                                                                                                                                                                                                                                              BCL2-associated X; BAX; nootropic; neuroprotective; antiparkinsonian; anticonvulsant; ophthalmological; antidiabetic; virucide; antisense therapy; BAX antagonist; BAX inhibitor; familial amylotrophic lateral sclerosis; Alzheimer's disease; Parkinson's disease; Hodgin's disease; cartilage-hair hyperplasia; diabetes-associated ocular disorder; scrapie infection; aberrant apoptosis; human; phosphorothioate; ss.
                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod base = OTHER
/note= "phosphorothicate linkages, and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                    Human BAX chimeric phosphorothicate oligonucleotide SEQ ID NO:26
                            ö
                            3; Indela
Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= a
/mod_base= OTHER
/note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Page 85; 139pp; English.
                                                        54 TCAGAGGAGTCTCTGCAC 71
                                                                                   N
                                                                                                                                                               B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-JUL-2001; 2001US-00908147.
 3.1%;
                                                                                     19 rchahadadccrrcracrc
                                                                                                                                                               ADA20853 standard; DNA; 20
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                                                                                                                                                                                                                       20-NOV-2003 (first entry)
               Local Similarity 83.3
hes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-239321/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zhang H, Watt AT;
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                            ADA20853;
                 Best Loca
Matches
                                                                                                                                  RESULT 406
                                                                                                                                                ADA20853
ID ADA2
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cc tissues comprising the expression of BAX protein in cells or tissues comprising contacting the cells or tissues with (1); and (3)

Cc treating an animal having a disease or condition associated with BAX

protein comprising administering to the animal (1) so that expression of

BAX protein is inhibited. (1) has nootropic, neuroprotective,

antiparkinsonian, anticonvulsant, ophthalmological, antidiabetic and

virucide activities, and can be used in antisense therapy, and as a BAX

attagonist. The antisense compounds (1) are useful for modulating the

expression of BAX protein, and for treating a disease or condition

associated with BAX protein, e.g. familial amylotrophic lateral

cartilage-hair hyperplasia, disease, parkinson's disease, Hodgkin's disease,

cartilage-hair hyperplasia, diabetes-associated ocular disorders or

scrapie infection, or a condition that arises from abbrrant apoptosis.

The compounds are useful as research reagents and in diagnostics. The

present sequence represents a human BAX chimeric phosphorothioate

oligonucleotide, which is used in an example from the present invention.

Sequence 20 BP; 1 A; 9 C; 7 G; 3 T; 0 U; 0 Other;

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Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
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ADA26274 standard; DNA; 20 BP 20-NOV-2003 (first entry) ADA26274; RESULT 407 ADA26274/C 

Zebrafish genomic DNA PCR primer #2.

Zebrafish, PCR; ss; hedgehog; neuronal cell; skeletogenesis; chrondrogenesis; osteogenesis; degenerative disorder; nervous system; neuronal cell death; neural cell; neuromuscular disorder; autonomic disorder; central nervous system disorder; anoxia; ischaemia; peripheral nervous system disorder; tachycardia; atrial cardiac arrhythmia; striated heart; stem cell development; digestive tract; liver; multiple sclerosis; primer.

Danio rerio.

US2003054437-A1.

20-MAR-2003

97US-00954771. 93US-00176427. 20-OCT-1997;

94US-00356060. 95US-00435093. 95US-00462386. 30-DEC-1993; 14-DEC-1994; 04-MAY-1995; 05-JUN-1995;

(INGH/) INGHAM P W. (MCMA/) MCMAHON A P. (TABI/) TABIN C J.

Tabin CJ; ingham PW, Mcmahon AP,

WPI; 2003-555377/52

Modulating growth, differentiation or survival of a cell, useful for treating a degenerative disorder of the nervous system characterized by neuronal cell death, comprises contacting the cell with a hedgehog polypeptide.

Example 4; Page 44; 121pp; English.

The invention relates to a method for modulating growth, differentiation or survival of a cell, comprising contacting the cell with a hedgehog polypeptide. The invention also relates to methods for inducing a cell to differentiate to a neuronal cell phenotype comprising a cell to differentiate to a neuronal cell phenotype comprising contacting a cell to carget tissue of a hedgehog polypeptide commission contacting a degenerative disorder of the nervous system characterised by neuronal cell death, comprising administering a hedgehog polypeptide causing prolonged survival of neural cells in the patient, relative to the absence of hedgehog propeptide are using prolonged survival of neural cells in the patient, relative to the absence of hedgehog treatment. The hedgehog polypeptides are using a degenerative disorder of the nervous system characterised by neuronal cell death, including content of the nervous system characterised by neurons of hedgens are disorders, including content and social sease, Parkinson's disease, multiple content and social demase resulting from anoxia, ischaemia or trauma and neuronal degeneration associated with a natural aging process. The neuronal degeneration associated with a natural aging process. The neuronal degeneration associated with a natural aging process. The chosen including disorders affecting innervation of smooth muscle and disorders affecting innervation of smooth muscle and peripheral nerve damage, for treating neoplastic or hyperplastic carriptement of stranformations and in controlling the development of stem cells crepairing central responsible for the heart, in nerve proscheses for repairing central responsible for the formation of the digestive tract, liver and other organs. This sequence represents a PCR primer used to amplify zebrafish 

Sequence 20 BP; 3 A; 6 C; 2 G; 1 T; 0 U; 8 Other;

Gaps . 0 3.1%; Score 13.2; DB 1; Length 20; 60.0%; Pred. No. 3.8e+02; rative 2; Mismatches 6; Indels Best Local Similarity 60.0 Matches 12, Conservative Query Match

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ACF57119 standard; DNA; 20 14-OCT-2003 (first entry) ACF57119; RESULT 408 ACF57119 PY X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B

BP.

Human sulfatase related probe SEQ ID NO:9.

Human; sulfatase; enzyme; cytostatic; neuroprotective; nootropic; antiparkinsonian; cerebroprotective; analgesic; cardiovascular; cardiant; antiansonian; antiarrhythmic; antianaemic; nephrotropic; uropathic; antialammatory; vasotropic; antiathmatic; gene therapy; cancer; CNS disorder; COPD; central nervous system disorder; cardiovascular disorder; asthma; haematological disorder; genitourinary disorder; chromosome X; Xp22.33; chronic obstructive pulmonary disease; probe; ss.

Homo sapiens. Synthetic.

WO2003057869-A1.

09-JAN-2003; 2003WO-EP000137 17-JUL-2003.

14-JAN-2002; 2002US-0347247P. 29-JUL-2002; 2002US-0398732P.

(FARB ) BAYER AG

Liou J;

rng.res

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The present invention describes a human sulfatase enzyme (I), which is located on chromosome X (more specifically to Xp22.33). (I) has cardiant, neuroprotective, antiparkinsonian, analgesic, cerebroprotective, cardiovascular, antiantantor, antianaemic, cerebroprotective, cardiovascular, antianthythmic, antianaemic, cerebroprotective, cardiovascular, antianthythmic, antianaemic, cerebroprotective, cardiovascular, antianthythmic, antianaemic, cordivities, and can be used in gene therapy. The sulfatase polymuclectide and polypeptide sequences can be used in diagnosing, preventing, candiorating or treating diseases associated with sulfatase dysfunction. They may also be used to identify test compounds that may act, for example, as activators or inhibitors at the enzyme's active site. The human sulfatase and its fragments are also useful in raising specific antibodies that can block the enzyme and effectively reduce its activity. The sulfatase can be used in the treatment of diseases such as cancer, a central nervous system (CNS) disorder (e.g. Alzheimer's disease, cordivators or parkinson's disease, stroke or neuropathic pain), a cardiovascular cordisorder (e.g. heart failure or arrhythmias), a haematological disorder (e.g. anaemia or thrombocytopaenia), a genitourinary disorder (e.g. renal cordinary disease (CDP)) or asthma. The present corporation obstructive pulmonary disease (CDP) or asthma. The present corporation or example from the present invention
                                                                                                                   New polynucleotide encoding a sulfatase polypeptide, useful for diagnosing, preventing or treating diseases associated with sulfatase dysfunction, e.g. cancer, asthma or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 20 BP; 4 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                         Example 16; Page 99; 124pp; English
WPI; 2003-577524/54.
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Score 13.2; DB 1; Length 20; Pred. No. 3.8e+02; 0; Mismatches 3; Indels ; 0 3.1%; Query Match
Best Local Similarity 83.3
Matches 15; Conservative ઠે

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Gaps ; 0

PKA regulatory subunit RII alpha inhibitory oligonucleotide ISIS102778. ACD44752 standard; DNA; 20 BP (first entry) 09-SEP-2003 ACD44752; 셤 

Human; ss; antisense therapy; infection; inflammation; tumour; protein kinase A regulatory subunit RII alpha. Synthetic

Homo sapiens

US6524854-B1 25-FEB-2003 11-SEP-2001; 2001US-00954560

(ISIS-) ISIS PHARM INC

11-SEP-2001; 2001US-00954560.

Monia BP,

WPI; 2003-511923/48.

New antisense compounds, useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha, and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha.

Example 15; Col 43-44; 35pp; English

The invention relates to antisense compounds targeted to nucleic acids encoding protein kinase A regulatory subunit RII alpha. The antisense compounds are useful for modulating the expression of protein kinase A (PKA) regulatory subunit RII alpha and for treating a disease or condition associated with expression of PKA regulatory subunit RII alpha. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human protein kinase A regulatory subunit RII alpha inhibitory oligomucleotide 

Seguence 20 BP; 3 A; 7 C; 9 G; 1 T; 0 U; 0 Other;

Gaps .; 0 Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

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320 CGTGCTGGCGGCGACGA 337 CATGCCGGCGGCGGA 18

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聁. ADB89866 standard; DNA; 20 04-DEC-2003 (first entry) ADB89866; RESULT 410 ADB89866/c

Antinsense oligonucleotide targeting human C3 component, ISIS139968. Human; 88; antisense; complement component C3; inflammation; septic shock; multiple organ fallure; hyperacute organ failure; autoimmune disorder; CNS inflammation; multiple sclerosis; atherosclerosis; tumour.

/mod\_base= OTHER /note= "Phosphorothioate backbone and all cytosines are -methyl cytosines" Location/Qualifiers Д Key modified\_base modified\_base Homo sapiens 

/\*tag= a /mod base= OTHER /note= "2'-methoxyethyl nucleotides" 16..20 /\*tag= c /mod\_base= CTHER /note= "2'-methoxyethyl nucleotides" modified\_base

US2003096775-A1

23-OCT-2001; 2001US-00001076 22-MAY-2003.

23-OCT-2001; 2001US-00001076.

SISI (-SISI)

Graham MJ,

WPI; 2003-606441/57. 

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The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid molecule encoding complement component C3. The compound specifically hybridises with the nucleic acid molecule encoding complement component C3, or specifically hybridises with at least an 8-nucleobase component C3, or specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding complement component C3. Also included are a composition comprising the component C3 in cells or tissues (comprising the expression of complement component C3 in cells or tissues (comprising complement C3 in cells or tissues) component C3 in cells or tissues with complement component C3 compounds are useful for inhibiting the expression of complement component C3 compounds are useful for inhibiting the expression of complement C3 in cells or tissues, or for treating an animal having a compound cited above so that compound can associated with complement component C3 in cells or tissues, or for treating an animal having a cutoinmune disorder (e.g. multiple solerosis), an infection, or atherosclerosis, inflammation, septic shock, multiple organ failure, cuseful as research reagents and diagnostics, in distinguishing functions of various members of a biological pathway, or for preventing or delaying complement component c3 in distinguishing functions inflammation or tumour formation. The present sequence is an entison or infection, inflammation or tumour formation. The present sequence is an entison or tumour formation. The present sequence is an entison or tumour formation. The present sequence is an entison or tumour formation. The present sequence is an entison or tumour formation. The present sequence is an entison or tumour formation. The present sequence is an entison or tumour formation. The present sequence is an entison or tumour formation. The present is the present is an entison or complement to the present is an entison or complement to the present is an entison or complement to the pre
New antisense oligonucleotides targeted to a nucleic acid molecule encoding complement component C3, useful for treating a disease or condition associated with complement component C3, e.g. autolmmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotide targeting human C3
                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 25; 72pp; English.
                                                                                                                                                                                                                                                  disorder or infection.
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ö Gaps .. 0 ch 1 Similarity 83.3%; Pred. No. 3.8e+02; 15; Conservative 0; Mismatches 3; Indels Sequence 20 BP; 3 A; 8 C; 6 G; 3 T; 0 U; 0 Other; Query Match Best Local Similarity Matches

321 GIGCIGGCGCGGACGAC 338 18 GIGCIGGAGGCCCACGAC 1 셤 ઠ

ADB68562 standard; DNA; 20 BP. 04-DEC-2003 ADB68562; RESULT 41 ADB68562/ 

(first entry)

homogeneous conjugate; hepatic; chronic viral hepatitis; cirrhosis; malaria; viral infection; protozoan; cancer; hepatocellular carcinoma; DNA oligonucleotide 9 targeted to Hepatitis C virus sequence.

Hepatitis C virus. HCC, HCV; ss.

402003067209-A2

14-AUG-2003.

21-JUN-2002; 2002WO-US019908

2-JUN-2001; 2001US-00888164

(CELL-) CELL WORKS INC. (UYJO ) UNIV JOHNS HOPKINS.

ΰ ຜ Deamond Duff R, Zhou Y, rs'o POP,

WPI; 2003-697456/66

New homogeneous prodrug conjugate containing hepatic ligand for delivery of pathogen-specific oligomer useful for treating liver infections or

Disclosure, Page 23; 107pp; English.

The invention relates to a novel homogeneous conjugate comprising a hepatic ligand, bifunctional linker and biologically stable oligomer that binds to a sequence in a hepatic virus or pathogen and is released from the conjugate by hydrolysis or reduction. The conjugate of the invention may be useful during the treatment of liver diseases including chronic viral hepatitis, cirrhosis, malaria, viral or protozoan infection and cancer, such as hepatocellular carcinoma (HCC). The current sequence is that of the DNA oligonucleotide 9 of the invention which is targeted to Hepatitis C virus (HCV) sequence. 

Sequence 20 BP; 2 A; 6 C; 6 G; 6 T; 0 U; 0 Other;

ö 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; ative 0; Mismatches 3; Indels Query Match
Best Local Similarity 85...
Best Local Si Conservative

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8

ADC71183 standard; DNA; 20 BP.

(first entry) 18-DEC-2003 ADC71183;

Nested PCR primer 2 (NP2) used in SSH to isolate 205P1B5 cDNA fragment.

205P1B5, prostate cancer; immune response; transgenic; knock out animal; cytostatic; immunogenic; vaccine; ss; SSH; suppressive subtractive hybridisation; PCR; primer; NP2.

Unidentified

WO2003020954-A2.

13-MAR-2003.

30-AUG-2002; 2002WO-US027760. 

31-AUG-2001; 2001US-0316664P.

(AGEN-) AGENSYS INC.

WPI; 2003-354484/33.

Jakobovits A; Hubert RS, Faris M, Raitano AB, Challita-Eid PM, ... volynucleotide designated 205PlB5, for diagnosing and treating prostate cancer, and as probes or primers for the amplification and/or detection of 205PlB5 genes.

Example 1; Page 60; 162pp; English.

This invention relates to a novel gene designated 205P1B5, and the encoded protein, which is aberrantly expressed in prostate cancer. Specifically, it refers to the two variants of 205P1B5 mapped to chromosome 8p21-8p12, namely 205P1B5v1 and 205P1B5v2 and fragments thereof that serve as useful diagnostic, prophylactic, prognostic, propressit cand, or therapeutic targets for prostate and other types of cancers. The present invention describes methods for the isolation of 205P1B5, for generating an immune response and for generating transgenic or knock out animals for

Sequence 20 BP; 3 A; 3 C; 7 G; 7 T; 0 U; 0 Other;

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Page 199

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the development and screening of therapeutically useful reagents. Furthermore, it refers to identifying proteins, small molecules or other agents that interact with 208PIBS, and can be used to identify pathways activated by 208PIBS. Accordingly, these are cytostatic and immunogenic compositions that are useful for the development of cancer vaccines. This suppressive subtractive is the nested PCR primer 2 (NP2) used for suppressive subtractive hybridisation (SSH) to isolate the 208PIBS cDNA fragment of the invention.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ; 0 3.1%; Score 13.2; DB 1; Length 20; 83.3%; Pred. No. 3.8e+02; arive 0; Mismatches 3; Indels 373 TCCTGGACCGCGACGACG 390 m 20 recresecescades Query Match
Best Local Similarity 83.35
Matches 15; Conservative ð 셤

RESULT 413

ADC16779 standard; DNA; 20 BP ADC16779; ADC16779/ 

18-DEC-2003 (first entry)

RT-PCR, PCR, primer, anti-retroviral, T-20, T-1249, S-Helix, env; gp41, anti-HIV; vaccine, albumin fusion protein; HIV fusion inhibiting peptide;

Forward RT-PCR primer to amplify HIV-1 RNA in order clone T-20.

ss; cyanovirin-N

Unidentified

WO2003066078-A1

14-AUG-2003.

07-FEB-2003; 2003WO-IB000434.

07-FEB-2002; 2002US-0355547P.

(AVET ) AVENTIS BEHRING GMBH. (DELZ ) DELTA BIOTECHNOLOGY LTD.

Hauser H, Weimer T, Sleep D;

WPI; 2003-731478/69.

New albumin fusion protein comprising a human immunodeficiency virus (HIV) fusion inhibiting peptide and an albumin having an albumin activity, useful for treating a disease or disorder, e.g. HIV infection.

Example 3; Page 58; 105pp; English.

This invention relates to novel albumin fusion proteins comprising a human immunodeficiency virus (HIV) fusion inhibiting peptide, which exhibit anti-retroviral activity. Specifically, it refers to inhibitory peptides including T-20, T-1249, 5-Helix or cyanovirin-N that bind the HIV env protein, or derivatives thereof such as the HIV gp41 protein. The Furthermore, the albumin activity has the ability to prolong the in vivo Fulfile of these HIV fusion inhibiting peptides. Accordingly, the present invention describes fusion proteins that neutralise HIV in a host or arising an immune response and also antibodies that inhibit viral infection of uninfected cells. In this way, a method exists to prevent, treat or ameliorate HIV infection and/ or a disease caused by HIV infection and/ or a disease caused by HIV infection and/ or a disease caused by HIV infection and or a disease caused by HIV infection and or a disease caused by HIV activity and can be used towards the production of a vaccine. This oligonucleotide sequence is the forward RT-PCR primer used to amplify the HIV-1 RNA in order to clone T-20, in an exemplification of the invention.

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Gape ö

Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels

119 CAAGTACGGCATGCTGGC 136

ò g

20

3 CAAGTATGTCATGCTGCC

ADD84533 standard; DNA; 20

RESULT 415

ADD84533

ADD84533;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a method of diagnosing endometriosis in a female suspected of having endometriosis comprising detecting the presence of a purified and isolated endometriotic haptoglobhr [BNDO-1) and its functional analogues from a patient sample. The presence of the endometriotic haptoglobin is indicative of endometriosis. The invention provides purified and isolated glycoprotein and biologically functional analogues having specific physical and functional characteristics. This sequence represents a primer used in the isolation of rat endometriotic haptoglobin ENDO-1 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diagnose of endometriosis in female involves detecting the presence of purified and isolated endometriotic haptoglobin and its functional
                                                                                                                                                                                                                                                                                            cytostatic; gynaecological; endometriosis; endometriotic haptoglobin;
ENDO-1; rat; PCR; primer; ss.
                               Gaps
                               ;
0
 Length 20;
                               Indels
                                                                                                                                                                                                                                                                Rat endometriotic haptoglobin ENDO-1 primer seq id 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
Match 3.1%; Score 13.2; DB 1; Local Similarity 83.3%; Pred. No. 3.8e+02; les 15; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; SEQ ID NO 7; 28pp; English.
                                                               172 ACTACGAGTCCAAGGCAC 189
                                                                                         18 ACTAGGATTCCAAGGCAC 1
                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94US-00328451.
98US-00044604.
                                                                                                                                                                                                                                                                                                                                                                                                                                           27-NOV-2002; 2002US-00306903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  analogs from patient sample.
                                                                                                                                                                        ADC78704 standard; DNA; 20
                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-802186/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (TIMM/) TIMMS K L.
                                                                                                                                                                                                                                                                                                                                                                              US2003166014-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-OCT-1994;
19-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                  Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Timms KL;
                                                                                                                                                                                                       ADC78704;
    Query Match
                                  Matches
                                                                                                                                             RESULT 414
                                                                                                                                                            ADC78704
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121P1F1; 121P1F1 modulation; human; chromosome 4q; cytostatic; gene therapy; vaccine; cancer; immune response; immunisation; primer; ss.
           121P1F1 gene nested primer (NP) 2 SEQ ID NO:721.
   29-JAN-2004 (first entry)
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The present invention describes a composition (I) comprising a substance that modulates the status of 121PIF1 (gene and encoded protein), or a molecule that is modulated by 121PIF1, where the status of a cell that expresses 121PIF1 is modulated. The human 121PIF1 gene maps to chromosome 4g. (I) has cytostatic activity, and can be used in gene therapy, and in vaccines. The composition (I) can be used for diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIF1, such as breast, colon, ovarian or lung cancer. The 121PIF1 gene or its fragment can be used to elicit a humoral or cellular immune response.

121PIF1 antibodies can be used in active or passive immunisation. 121PIF1 polynucleotides are useful as probes and primers for the amplification or detection of 121PIF1 genes, as coding sequences for the amplification or inhibiting the expression of 121PIF1 genes. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Composition comprising a substance that modulates the status of 121PIF1, useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIF1, such as breast, colon, ovarian or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Challita-Eid PM, Hubert RS, Raitano AB, Faris M, Afar DEH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 71; 285pp; English.
                                                                                                                                                                                                                                                                         28-FEB-2002; 2002WO-US006242.
                                                                                                                                                                                                                                                                                                                                                                  05-MAR-2001; 2001US-00799250.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AGEN-) AGENSYS INC.
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                                                                                   WO200295009-A2.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lung cancer
                                                                                                                                                                                 28-NOV-2002
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This invention relates to a novel composition which comprises a substance that modulates the status of a novel protein (1612/2108) and its variants braving a sequence of 875 amino acids provided in the specification. The protein of the invention is over-expressed in certain cancers. The compounds of the invention may have cytostatic activity and the sequence of the IEDP2NOB protein, and the gene which encodes it, may be useful for gene therapy or the development of a vaccine. The composition and methods of the invention are useful in diagnosing, preventing and treating ancourt. The present sequence is that of FCR primer which was used for amplification of a region of the gene encoding the human 161P2F10B protein during the exemplification of the invention.

A composition for diagnosing, preventing and treating cancer (e.g. prostatic, renal or uterine cancer) comprises 161P2F10B polynucleotides and polypeptides.

Example 1; SEQ ID NO 36; 135pp; English.

Morrison KJM;

Ge ₩

A, Raitano AB, Faris M, Hubert RS, K, Challita-Eid PM;

(AGEN-) AGENSYS INC

WPI; 2003-441560/41.

Jakobovits A, Morrison RK,

07-NOV-2002; 2002WO-US036002. 07-NOV-2001; 2001US-00005480 31-JAN-2002; 2002US-00062109

WO2003040340-A2

15-MAY-2003

Homo sapiens.

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                                                                                                                                                                                                                                                      193PIE1B; tissue specific expression, cancer, cytostatic, gene therapy, cancer, human, PCR, RT-PCR, reverse transcription PCR, primer, ss.
                                               Gaps
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                    Query Match 3.1%; Score 13.2; DB 1; Length 20; Best Local Similarity 83.3%; Pred. No. 3.8e+02; Matches 15; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Hubert RS,
Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                               Human protein 193P1E1B-related PCR primer SeqID59.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Faris M,
                                                                        373 TCCTGGACCGCGACGACG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                         Challita-Eid PM,
                                                                                      20 TCCTCGGCCGCGACCACG 3
                                                                                                                                                           ВÞ.
                                                                                                                                                                                                                                                                                                                                                                  06-DEC-2002; 2002WO-US039274.
                                                                                                                                                                                                                                                                                                                                                                                          07-DEC-2001; 2001US-00013312.
                                                                                                                                                           ADD96944 Btandard; DNA; 20
                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  (AGEN-) AGENSYS INC.
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                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                         29-JAN-2004
                                                                                                                                                                                                                                                                                                                                           19-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Raitano AB,
                                                                                                                                                                                  ADD96944;
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                                                                                                                                    RESULT 417
ADD96944/c
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Gaps

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373 TCCTGGACCGCGACGACG 390

à g

50

161P2F10B; cancer; cytostatic; gene therapy; vaccine; PCR; primer; ss;

Human 161P2F10B protein-related PCR primer SeqID36.

(first entry)

BP.

ADE65924 standard; DNA; 20

ADE65924;

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This invention relates to novel composition comprising a substance that modulates the status of a 433 residue protein, given in the specification with the DNA sequence encoding it, or a molecule that is modulated by the protein. The novel protein 193P121B exhibits tissue specific expression in normal adult tissue and is aberrantly expressed in certain cancers. Compositions which modulate the 193P121B protein may have cytogratic activity and the DNA sequence which encodes protein 193P121B may be useful in gene therapy. The composition of the invention may be useful primer which was used for the amplification of human 193P121B gene DNA during the exemplification of the invention.
                                                                                                                                                                                          New composition comprising 193P1E18-related protein, useful for preventing or treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID NO 59; 260pp; English.
                                                                                               WPI; 2003-532905/50.
Jakobovits A;
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Score 13.2; DB 1; Length 20;
Pred. No. 3.8e+02;
); Mismatches 3; Indels
                                            373 TCCTGGACCGCGACGACG 390
                      ;
0
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 3.1%;
                                                                 rccrcecceceaceace
                       Conservative
          Local Similarity
Les 15; Conser
                                                                 20
 Query Match
                      Matches
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Wild-type capture oligonucleotide #13 AAF95086 standard; DNA; 16 BP (first entry) 23-MAY-2001 AAF95086; 

Tubercle bacillus, drug sensitivity, drug resistance, rifampicin, streptomycin, kanamycin, isoniazid, ethambutol, rpoB gene, rrs gene, rpsL gene, inhA gene, katG gene, embB gene, probe, PCR primer, ss.

Mycobacterium tuberculosis

EP1076099-A2

14-FEB-2001.

02-AUG-2000; 2000EP-00306563

99JP-00220357. 03-AUG-1999;

(NISM ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

Takenishi S; Suzuki Y, Nishida M,

WPI; 2001-246696/26.

New oligonucleotides, nucleic acid probes and primers are useful for differentiating drug-resistance and determining infection with tubercle bacilli.

Claim 21; Page 40; 114pp; English.

The present invention relates to oligonucleotides based on nucleotide sequences obtained from both wild-type tubercle bacilli (wrTB) that are succeptible to a drug and mutant-type tubercle bacilli (mrTB) that are resistant to a drug. The drugs used in the present invention are

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rifampicin (RPP), streptomycin (SM), kanamycin (KM), isoniazid (INH) and ethambutol (EB). The trops gene is responsible for resistance to RPP; the responsible for resistance to SM and KM, the rpst gene is responsible for resistance to SM, the inhA gene is responsible for resistance to INH; the katG gene is responsible for resistance to INH; and the embB gene is responsible for resistance to INH; and the embB gene is responsible for resistance to INH; invention also relates to nucleic acid probes having part of a nucleotide primers used to generate the probes. The present sequence is an oligonucleotide of the present invention. The oligonucleotides of the present invention. The oligonucleotides of the spread to enable the differentiation of drug resistance and the determination of infection with tubercle bacilli simultaneously
             888888888888888888
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Sequence 16 BP; 3 A; 4 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ô Length 16; 0; Indels 3.1%; Score 13; DB 1; Le 100.0%; Pred. No. 2.6e+02; :ive 0; Mismatches 0; Local Similarity 100. les 13, Conservative Query Match Best Loc

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Hammerhead ribozyme substrate #323. ВP AAF02028 standard; DNA; 17 16-FEB-2001 AAF02028;

RESULT 419

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Gaps

; 0

Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss. Homo sapiens.

11-APR-2000; 2000WO-US009721. WO200061729-A2. 19-OCT-2000. 

99US-0129390P (RIBO-) RIBOZYME PHARM INC. 12-APR-1999;

Mcswiggen J; Blatt L, Zwick M, Pavco P,

WPI; 2000-647423/62.

ssor genes, factor protein, Enzymatic and antisense nucleic acid inhibition of repressor useful for producing e.g. granulocyte colony stimulating fact interferon alpha and erythropoietin.

Claim 37; Page 63; 164pp; English.

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRP-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha

Sequence 17 BP; 1 A; 9 C; 6 G; 1 T; 0 U; 0 Other;

Length 17; 3.1%; Score 13; DB 1; Lt 100.0%; Pred. No. 2.9e+02; Query Match Best Local Similarity Length 17;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                            Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene;
Rho GTPase; signal transduction; gene expression; cancer; vaccine;
gene therapy; transgenic; ss.
 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; SEQ ID NO 1749; 60pp + Sequence Listing; English.
 Indels
                                                                                                                                                                                                                                                                Human POSHL1 scanning oligonucleotide SEQ ID NO 1749.
;
 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-US000669.
                                                                                                                                                           ABV91036 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JAN-2002; 2002EP-00001165.
                                  248 CCCGGGCTCGGCC 260
                                                                                                                                                                                                                                23-DEC-2002 (first entry)
                                                          1 CCCGGGCTCGGCC 13
 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                              EP1239051-A2.
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shannon M;
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   Matches
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Sequence 17 BP; 2 A; 7 C; 7 G; 1 T; 0 U; 0 Other;

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (SI, ABB3939), a sequence having 65$ sequence identity to (SI), (SI) having 95$ deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene, foncedent for that functions as an adaptor protein that interacts with Rho family small Grasses as well as downstream components of the signal transduction pathway. (1) is useful of or identifying a specific binding partner. (7) and nucleic acids (11) for oused by altered expression of human POSHL1 including diagnosing and caused by altered expression of human POSHL1 including diagnosing and causeful for measuring and for surveying gene expression and creating are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                            Human; POSH1 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 1752; 60pp + Sequence Listing; English.
Query Match 3.1%; Score 13; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                           Human POSHL1 scanning oligonucleotide SEQ ID NO 1752.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JAN-2001; 2001MO-US000664.
30-JAN-2001; 2001MO-US000664.
30-JAN-2001; 2001MO-US000665.
30-JAN-2001; 2001MO-US000666.
30-JAN-2001; 2001MO-US000668.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001MO-US000669.
30-JAN-2001; 2001WG-US000670.
23-MAY-2001; 2001US-00864761.
                                                                                                                                                                                                                 ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JAN-2002; 2002EP-00001165
                                                                                                                                                                                                                 ABV91039 standard; DNA; 17
                                                                             338 CCAGGGCCGGCTG 350
                                                                                                                                                                                                                                                                                          23-DEC-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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ABV91039/
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RESULT 42:

ABV91038/1

AXC ABV9:

XX ABV9:

DDT 23-Di

DX 23-Di

DX BRD 6

XX Humai

XX Humai

XX Homo

YX ABV9:

YX A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                  Gaps
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                                                                                                                     DB 1; Length 17; 2.9e+02;
                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human POSHL1 scanning oligonucleotide SEQ ID NO 1750.
                                                                 Sequence 17 BP; 2 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                            3.1%; Score 13; DB 100.0%; Pred. No. 2.9 tive 0; Mismatches
Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001, 2001WO-US000663.
30-JAN-2001, 2001WO-US000664.
30-JAN-2001, 2001WO-US000666.
30-JAN-2001, 2001WO-US000666.
30-JAN-2001, 2001WO-US000667.
30-JAN-2001, 2001WO-US000669.
30-JAN-2001, 2001WO-US000669.
30-JAN-2001, 2001WO-US000670.
23-MAY-2001, 2001US-00864761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV91037 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                             338 CCAGGGCCGGCTG 350
                                                                                                           Ouery Match
Best Local Similarity 100.0
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                          CCAGGGCCGGCTG 1
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ABV91037/c
ID ABV9103
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present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to Derwent by the Buropean Patent Office
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Rho GTPase; signal transduction; gene expression; cancer; vaccine;
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                                                                                                                                                                                                                                     Length 17;
                                                                                                                                                                                                                                                                                                    0; Indels
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                                                                                                                                                                                                                                  Score 13; DB 1; Le
Pred. No. 2.9e+02;
0; Mismatches 0;
                                                                                                                                                                       Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;
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100.0%; Pred
0; M
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US0006667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-US006670.
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                                                                                                                                                                                                                                                                                                                                                                   338 CCAGGGCCGCTG 350
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                                                                                                                                                                                                                                                                     Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                   15 CCAGGCCGGCTG 3
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                                                                                                                                                                                                                                     Query Match
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useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence did not form part of the Derwent by the Buropean Patent Office

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Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

0; Gaps Query Match 3.1%; Score 13; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 0; Indels 338 CCAGGGCCGGCTG 350 ઠે

ACC65163 standard; DNA; 17 BP. RESULT 424 ACC65163

ACC65163;

01-JUL-2003 (first entry)

Murine oligonucleotide associated with tumour supression, SEQ ID 2410.

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.

Mus musculus.

WO2003025176-A2.

27-MAR-2003.

17-SEP-2002; 2002WO-IB004210.

17-SEP-2001; 2001FR-00011979.

(MOLE-) MOLECULAR ENGINES LAB

Tuijnder M; Telerman A, Amson R,

New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells. WPI; 2003-333167/31.

Disclosure; Page 312; 738pp; French.

The present invention relates to murine oligonucleotides (ACC62754-ACC68066), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia 

Sequence 17 BP; 3 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

Gaps . 3.1%; Score 13; DB 1; Length 17; 100.0%; Pred. No. 2.9e+02; ive 0; Mismatches 0; Indels Best Local Similarity 100. Matches 13; Conservative

2 Arceeaaecrec 14

CC

AAA38383 standard; DNA; 18

AAA38383;

21-AUG-2000 (first entry)

Ets-2; human; transcription factor; chromosome 21q22.3; cancer; invasion; metastasis; skeletal abnormality; Down's syndrome; expression inhibition; phosphorothioate; antisense; ss.

Human Ets-2 phosphorothioate antisense oligonucleotide, SEQ ID NO:42.

Homo sapiens

US6054316-A.

25-APR-2000.

99US-00344579 25-JUN-1999; 99US-00344579. 25-JUN-1999;

(ISIS-) ISIS PHARM INC.

Baker BF, Cowsert LM;

WPI; 2000-338495/29.

Antisense compound, 8-30 nucleobases in length, inhibiting the expression Ets-2 is useful for treating cancer and detecting Ets-2 expression.

Claim 3; Col 40; 31pp; English.

Sequences AAA38349-A38388 represent antisense oligonuclectides targetted to the human Ets-2 gene, which inhibit its expression. The antisense oligonuclectides were designed to target different regions of the human Ets-2 RNA, and were analysed for their effect on Ets-2 RNA levels by Ets-2 RNA levels by Camilitative real-time FCR. The Ets-domain transcription factors are a family of proteins which are involved in controlling key callular events contains a proliferation, differentiation and development. The Ets domain is a DNA-binding domain shared by all members of this family. Through this cat a GCA consentus sequence, thereby acting as either repressors or atta GCA consentus sequence, thereby acting as either repressors or activators of the gene. All but one Ets family protein bind to DNA as a conomer. Ets-2 has been implicated in the regulation of callular corrections associated with malignancies. Ets-2 has been found to be translocations associated with malignancies. Ets-2 has been found to be translocations associated with malignancies. Ets-2 has been found to be translocations associated with malignancies. Ets-2 has been found to be upregulated in several cancers, including lymphoblastic leukaemia. It may calso play a role in the cancer phenotype, as it activates the urokinase of plasminogen activator (upp) promoter and the promoters of continuation. High levels of upp and metalloproteinases are associated with tumour invasion and metatasis in breast cancers. As the Ets-2 gene is located on chromosome 21, which is triplicated in Down's syndrome, it is also thought to be responsible for the skeletal abnormalities present to this condition. The artisense of conditions associated with Ets-2 expression, especially cancer 

Sequence 18 BP; 3 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

.. 0 Query Match
3.1%; Score 13; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 13; Conservative 0; Mismatches 0; Indels

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368 4

ò d ABK33430;

RESULT 426

ABK33430

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AAV46388-V46391 are PCR primers used in the isolation of a perchloroethane dehalogenase (PCE-DH) isolated from Dehalospirillum multivorans. This protein is used in a process for microbiological purfiication of water conteminated with chlorinated ethylenes and/or chlorinated propylenes. The process involves adding an electron donor a passing the water through a bioreactor containing a syntroppic mixed culture immobilised on a support, where the culture comprises at least one dehalogenating bacterium and at least one hydrogen-producing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Microbiological purification of water contaminated with chlorinated olefin(8) - using combination of dehalogenating and hydrogen-producing
                                                                    Perchloroethane dehalogenase; PCE-DH; microbiological purification; water contamination; chlorinated ethylene; propylene; electron donor; bioreactor; dehalogenating bacterium; anaerobic microorganism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13; DB 1; Length 20;
Pred. No. 4.1e+02;
1; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 4 A; 1 C; 8 G; 3 T; 0 U; 4 Other;
                                   D. multivorans PCE-Dehalogenase PCR primer #3.
                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 16; 27pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  one dehalogenating bacterium and strictly anaerobic microorganism
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/mod_base= i
/note= "inosine"
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/mod_base= i
/note= "inosine"
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/mod_base= i
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                                                                                                                                                                          Synthetic.
Sulfurospirillum multivorans.
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Best Local Similarity 72.2
Matches 13; Conservative
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modified_base
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Eisenbeis M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-MAR-1997;
18-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-SEP-1998,
                                                                                                                                 PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP864542-A2
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AAX21359/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method for detecting non-responders to anti-tumour necrosis factor (TNF) therapy. The method involves testing an individual for homozygosity for at least one single nucleotide polymorphism (SNP) in the gene coding for TNF receptor II, which is located on chromosome 1p36. Two novel SNPs, one in exon 2 (position 168 A/G) and one in exon 6 (position 5NPs, one in exon 2 (position 168 Met196Arg respectively, are also described. The method of the invention is useful for detecting non-responders to anti-TNF therapy such as infliximab therapy, or therapy of Crohn's disease. The genes containing the 2 novel polymorphisms are useful for diagnostic purposes in inflammatory, malignant or other chronic diseases. ABK33417-ABK33440 represent PCR primers used to amplify different regions of the human TNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting non-responders to anti-human necrosis factor therapy, comprises testing an individual for homozygosity for a single nucleotide polymorphism in the gene coding for the tumor necrosis factor receptor
                                                                                                                                                                                                                                                                                             Human; anti-tumour necrosis factor receptor II; TNF receptor II; chromosome 1p36; infliximab therapy; Crohn's disease; malignant disorder; inflammatory disorder; chronic disease; receptor; PCR; primer; ss.
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                                                                                                                                                                                                                                                           Human TNF receptor II gene exon 4 PCR primer #2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 4; 45pp; English.
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                                                                                                                                         ABK33430 standard; DNA; 19 BP
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      CACTITCCIGGAC 380
                                                                                                                                                                                                                    23-APR-2002 (first entry)
                           CACTTTCCTGGAC 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schreiber S,
                                                                                                                                                                                                                                                                                                                                                                                                                     EP1172444-A1.
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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AAV46390;

AAV46390 ID AAV4 XX AC AAV4 XX

RESULT 427

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Query Match

Matches

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This sequence represents a preferred antisense oligomucleotide targetted against the gene encoding human microtubule-associated protein 4 (MAP4). Inhibition of MAP4 expression was measured by determination of MAP4 mRNA levels in a variety of cell lines via real-time quantitative PCR. The cell lines used included the bladder carcinoma cell line PCR. The numan lung carcinoma cell line A549, human neonatal dermal fibroblasts and human embryonic keratinocytes. Microtubule-associated proteins comprise a group of proteins that mediate microtubule assembly and function which is required for cytoskeletal integrity. MAP4 is a member of the non-neuronal strabilising the microtubule lattice. MAP4 expression, and is therefore elevated in cells with mutant p53 oncogene expression, and is therefore linked to cancer chemcherapeutic drug sensitivity. These antisense concerning animals, particularly mamnas, having or being prone to a disease or condition associated with the expression of MAP4. The oligomucleotides are also useful for research and diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense oligonucleotides for inhibiting microtubule-associated protein 4 expression, useful in treating disorders associated with microtubule protein expression.
                                                                                                                              methoxyethyl (2'-MOE) nucleotides"
note= "2' methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence of PCR primer HCG-R2.
                                                                                                                   OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Col 39; 39pp; English.
                                                                                                                                                                                          mSc
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/mod_base= m5c
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                                                           m<sub>5</sub>C
                                 /mod_base= m.
16. .20
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                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                 *tag=
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                       modified base
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                                                                                                                                                                                                                                                                                                                                                             39-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                09-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61
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AAZ99376/c
     g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primers AAX21358-X21359 were used to PCR amplify a fragment of the 17DE1 locus sequence as a control sequence for analysis of BAI gene expression in blots. The BAI genes (see AAX21355-X2137) are expressed specifically in the brain and play an important role in cancer formation in the brain. The BAI proteins can be used in drug compositions to diagnose, prevent or
                                                                                                                                                                                                                                                                                                                                                                                                                                        New human BAI gene - is expressed in brain plays important role in cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Microtubule associated protein 4; MAD4; real-time quantitative PCR; expression; microtubule; assembly; function; cytoskeleton; structural; dynamic; stabilisation; lattice; overexpression; p53; oncogene; cancer; chemotherapy; tumour; drug sensitivity; antisense; therapy; hybridisation; inhibition; research; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                             Human; BAII; brain; cancer; drug; diagnosis; prevention; treatment;
primer; PCR; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human microtubule-associated protein 4 (MAP4) antisense oligo #37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 3.1%; Score 13; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 4.1e+02; les 13; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 2 A; 8 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; Page 16; 62pp; Japanese.
                                                               Prime ElA for 17DE1 locus sequence.
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/*tag= a
/mod_base= OTHER
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/*tag= b
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                           97JP-00176485.
                                                                                                                                                                                                                                                                                                                             97JP-00150460,
                                                                                                                                                                                                                                                                                                                                                              (SAKA ) OTSUKA PHARM CO LTD.
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                           21-MAY-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-183823/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treat such cancers
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modified_base
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                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                    JP11032766-A
                                                                                                                                                                                                                                                                                                                               23-MAY-1997;
                                                                                                                                                                                                                                                                                           16-JUN-1997;
                                                                                                                                                                                                                                                   09-FEB-1999.
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                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                formation
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WO200009734-A2.
                                                                                                                                          Unidentified.
                                      Mitchell LG,
                                                                                                                         03-JUL-2000
                       12-AUG-1999;
                           13-AUG-1998;
23-SEP-1998;
          Unidentified
                  24-FEB-2000
                                                                                                                     AAZ99396;
                                                                                         Query Match
                                                                                                              AAZ99396
                                                                                                            RESULT
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Thymosin-beta-10-like protein; ephrin type-A receptor 8-like protein; 85; proteoglycan-like protein; fibromodulin; fibromectin; thymic immune cell; spermatogenesis; male infertility; neoplasia; red blood cell; platelet; small cell lung cancer; GPI-anchored ephrin-A ligand; prostate cancer; neurological disorder; cardiac disorder; vascular disorder; orthopaedic; inflammatory disease; rheumatoid archritis; connective tissue; congenital muscular dystrophy; chemotherapy; immunotherapy; PCR primer; BC 2.7.1.112.
                                                                                                                                                                                                                                                                                                                                   The specification describes a pre-trans-splicing molecule (PTM) which contains one or more target binding domains, a 3' splice region comprising a branch point, a pyrimidine tract and a 3' splice acceptor site, a spacer region separating the mRNA splice region from the target binding domain, and a nucleotide sequence to be trans-spliced. The method is used for the in vivo production of a trans-spliced molecule in a subset of cells. The PTM is used for producing chimeric mRNA molecule by contacting it with target pre mRNA which is useful for gene regulation, gene repair and targeted cell death particularly repair of cystic fibrosis trans-membrane regulator gene. The present primer was used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Forward PCR primer for analysis of ephrin type-A receptor 8-like protein.
                                                                                                                                                                                                     Novel pre-trans-splicing molecules for use in gene regulation, gene repair and targeted cell death particularly gene repair of cystic fibrosis trans-membrane regulator gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cch 3.1%; Score 13; DB 1; Length 20; al Similarity 100.0%; Pred. No. 4.1e+02; 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                   Example 7; Page 42; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       test a lacZ trans-splicing model
                                                                                                               Garcia-Blanco MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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18-OCT-1999; 99US-0159992P.
22-OCT-1999; 99US-0160952P.
12-OCT-2000; 2000US-00159805.
98US-00133717.
98US-00158863.
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                                                                  (INTR-) INTRONN HOLDINGS LLC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      255 TCGGCCACGGTGC 267
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                                                                                                                                                             WPI; 2000-224360/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                                                                                 Mitchell LG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-SEP-2001
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  13-AUG-1998;
23-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes a pre-trans-splicing molecule (PTM) which contains one or more target binding domains, a 3' splice region comprising a branch point, a pyrimidine tract and a 3' splice region first site, a spacer region separating the mRNA splice region from the target binding domain, and a nucleotide sequence to be trans-spliced. The method is used for the in vivo production of a trans-spliced molecule in a subset of cells. The PTM is used for producing chimeric mRNA molecule by contacting it with target pre mRNA which is useful for gene regulation, gene repair and targeted cell death particularly repair of cystic fibrosis trans-membrane regulator gene. The present primer was used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel pre-trans-splicing molecules for use in gene regulation, gene repair and targeted cell death particularly gene repair of cystic fibrosis trans-membrane regulator gene.
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                        Pre-mRNA molecule, gene repair; pre-trans-splicing molecule, gene regulation; targeted cell death; cystic fibrosis trans-membrane regulator gene; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer HCG-R2 used to test a lacZ trans-splicing model,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.1%; Score 13; DB 1; Length 20;
100.0%; Pred. No. 4.1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 32; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                         Garcia-Blanco MA;
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98US-00158863
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ99396 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-224360/19.
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AAD38163 standard; DNA; 20 BP. 10-SEP-2002 (first entry) AAD38163; Вb

Membrane bound protein; secreted NOV protein; spermatogenesis; neoplasia; male infertility; angiogenesis; vascular pathology; orthopaedic disorder; inflammatory disease; congenital muscular distropy; muscular disorder; rheumatorid arthritis; fixed deformity; dysprothrombinaemia; oancer; arthrogryposis; hypoprothrombinaemia; hypokalaemic period paralysis; Smith-Lemil-Opitz syndrome; carcinoid tumour; centrocytic lymphoma; hypoprarathyroidism; Leigh syndrome; cervical carcinoma; leuksemia; macular dystrophy; vielliform type; McArdle disease; Meckel syndrome; parathyroid adenomatosis 1; multiple myeloma; hyperparathyroidism; parathyroid adenomatosis 1; prolactinoma; digenic retinitis pigmentosa; somatotrophinoma; neovascular inflammatory vitreoretinopathy; arthritis; carcinoid syndrome; atopy; tendonitis; gene therapy; vaccine; PCR; primer; ss. 

Inidentified

L8-APR-2002.

10-OCT-2001; 2001WO-US031498

The sequence represents a PCR primer used in expression analysis of ephrin type-A receptor 8-like protein (NOV1). Thymosin-beta-10-like protein (NOV1), ephrin type-A receptor 8-like protein and proteoglycan-common to the diagnosis, treatment and proteins (NOV3) may be used in the diagnosis, treatment and proteins (NOV3) may be used in the diagnosis, treatment and proteins (NOV3) may be used in the diagnosis, treatment and chirchnectin. The polypetides of the invention are useful in screening for agents that modulate their activity, and in determining predispositions to disorders. NOV1 is useful for treating conditions involving development, differentiation, and activation of thymic immune cells, in pathologies related to spermatogenesis and male infertility, diagnosis of neophasias, in diseases or pathologies of red blood cells or platelets, in detection of small cell lung cancer. NOV1 nucleic acids can be combined in chemo-immunotherapeutical anti-cancer treatments. NOV2 is useful for detecting cells expressing GPI-anchored ephrin-A ligands, as a marker for prostate cancer, and in treating cells acids and proteins are useful for treating orthopasedic disorders and/or injuries, and inflammatory diseases of connective tissues e.g. rheumatoid arthritis, congenital muscular dispendents. New isolated polypeptides, NOV 1-3, having identity to thymosin-beta-10, ephrin type-A receptor 8 and proteoglycans, and polymucheotides, useful for treating male infertility, neurological or cardiac disease or rheumatoid arthritis. Example 1; Page 83; 102pp; English.

0; Gaps Match 3.1%; Score 13; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 4.1e+02; tes 13; Conservative 0; Mismatches 0; Indels Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other; Query Match 

7 GAGTGAAACTGCG 19

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NOV2 cDNA specific forward PCR primer.

40200230979-A2.

12-OCT-2000; 2000US-00689486. 13-OCT-2000; 2000US-00687276. 09-OCT-2001; 2001US-00973424. 

(CURA-) CURAGEN CORP

ä Bandaru Prayaga SK, Taupier RJ,

WPI; 2002-454545/48.

Novel membrane bound and secreted NOV polypeptides, for treating, diagnosing and preventing male infertility, neurological, cardiac and arthritis.

Example 1; Page 118; 180pp; English.

The present invention relates to novel membrane bound and secreted NOV proteins and polynuclectides encoding such proteins. Sequences of the proteins and polynuclectides encoding such proteins. Sequences of the invention are useful for treating a medicament for treating a syndrome associated with human disease. They are useful for determining the associated with human disease. They are useful for determining the presence of or predisposition to lung cancer. NOVI compounds are useful for development, differentiation and activation of thymic immune cells, pathologies related to spermatogenesis and male infertility, diagnosis of several human neoplasias and diseases or pathologies of cells in blood circulation such as red blood cells and platelets. NOVI nucleic acids are useful for detecting specific cell types and as specific marker for cancers in tissues. NOV2 and NOV4 compounds are useful for treating system and angiogenesis and for treating cardiac and vascular pathologies. NOV3 and NOV5 compounds are useful for treating vascular pathologies. NOV3 and NOV5 compounds are useful for treating vascular pathologies. NOV3 and NOV5 compounds congenital muscular dystrophies, various muscular disorders indicated congenital muscular dystrophies, various muscular disorders, fixed for treating atopy, dysprothrombinaemia, hypoprothrombinaemia, type I and Correcting atopy, dysprothrombinaemia, hypoprothrombinaemia, type I and Cype, macular dystrophy, vitelliform type, McArdle disease, type 2 McCkellyphone, multiple endocrine neoplasial, multiple mysolome, multiple endocrine neoplasial, multiple mysolome, somatorrophinome, cype, macular dystrophy, vitelliform type, McArdle disease, type 2 McKellyphone, multiple endocrine neoplasial, multiple mysolome, somatorrophinome, sequence is specific por primer. This sequence is used in the exemplification of the invention

Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

the invention

ö Query Match
3.1%; Score 13; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 13; Conservative 0; Mismatches 0; Indels

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(first entry) 02-OCT-2002 ABQ73441;

ABO73441 standard; DNA; 20 BP.

RESULT 434 ABQ73441/

Human beta-chronic gonadotropin (HCG) RT-PCR primer HCG-R2.

Pre-trans-splicing molecule; PTM; spliceosome; cytostatic; gene therapy immunosuppressive; antimicrobial; gene regulation; gene repair; cancer; targeted cell death; genetic disorder; infectious disorder;

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Wed Apr 21 12:58:21 2004
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Novel cell having pre-trans-splicing molecules with target binding domains that target binding of PTM to pre-mRNA, 3' or 5' splice region, spacer region, nucleotide sequence to be trans-spliced to target-pre-mRNA.
                                                                                                                                          Garcia-Blanco MA, Baker CC, Puttaraju M;
Chao H;
autoimmune disease; proliferative disorder; PCR primer;
                                                                                                                                                                                                                      Example; Page 43; 229pp; English.
                                                                               08-JAN-2001; 2001US-00756095.

08-JAN-2001; 2001US-00756096.

08-JAN-2001; 2001US-00756097.

20-APR-2001; 2001US-00841895.
                                                                 08-JAN-2002; 2002WO-US000416
                                                                                                                             (INTR-) INTRONN INC.
                                                                                                                                                                  WPI; 2002-566693/60.
                                    WO200253581-A2.
                                                                                                                                           Mitchell LG,
Mansfield GS,
             Homo sapiens.
Synthetic.
                                                   11-JUL-2002
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The present invention describes a cell (I) comprising pre-trans-splicing molecules (PTMs) (II) which have one or more target binding demains (IIa) content and 3'splice region (IIb) that concludes branch point pyrimidine tract and 3'splice acceptor site, or 5' includes branch point pyrimidine tract and 3'splice acceptor site, or 5' splice site (IIC), spacer region (IId) that separates RNA splice site (IIC) spacer region (IId) that separates RNA splice site (IIC) and (IIE); or (B) (IIC) (IId) and (IIE) that comprises (II) be trans-comprising: (A) (ID) and (IIE); or (B) (IIC), (IId) and (IIE). The cell comprises (II) comprises (III) and (IIE), or (B) (IIC), (III) and (IIE), or (IIC), (III) and (IIE), or (IIC) (III) and (IIE), or (IIC) (III) and (IIE), or (IIC), (III) and (IIE), or (IIC) (IIC) and (II

Query Match

3.1%; Score 13; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 13; Conservative 0; Mismatches 0; Indels Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;

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RESULT 435 ABQ73457/c ID ABQ73457 standard; DNA; 20 XX

(first entry) 02-OCT-2002 ABQ73457; 

rng.res

Human beta-chronic gonadotropin (HCG) related PCR primer HCG-R2

Pre-trans-splicing molecule, PTM; spliceosome; cytostatic; gene immunosuppressive; antimidrobbial; gene regulation; gene repair; targeted cell death; genetic disorder; infectious disorder; autoimmune disease; proliferative disorder; FCR primer; 89.

Homo sapiens Synthetic.

WO200253581-A2.

11-JUL-2002.

08-JAN-2002; 2002WO-US000416

08-JAN-2001; 2001US-00756095. 08-JAN-2001; 2001US-00756096. 08-JAN-2001; 2001US-00756097. 20-ARR-2001; 2001US-00941492.

(INTR-) INTRONN INC.

Novel cell having pre-trans-splicing molecules with target binding domains that target binding of PTM to pre-mRNA, 3' or 5' splice region, spacer region, nucleotide sequence to be trans-spliced to target-pre-Garcia-Blanco MA, Baker CC, Puttaraju M; Chao H; WPI; 2002-566693/60. Mitchell LG, Mansfield GS,

Example; Page 53; 229pp; English.

The present invention describes a cell (I) comprising pre-trans-splicing molecules (PTMs) (II) which have one or more target binding domains (IIa) that target binding of PTM to pre-mRNA, 3' splice region (IIb) that circledes branch point pyrimidine tract and 3'splice acceptor site, or 5' splice site (IIc), spacer region (IId) that separates RNA splice site from target binding domain, and mucleotide sequence to (IIe) be transcomprising: (A) (IIb) and (IIe), the cell comprises (II) either comprising: (A) (IIb) and antimicrobial activities, and can be used in gene therapy. (II) comprising one or more (preferably two or more) (IIa) and (IIe), or (II) (IId) and (IIe), or (II) comprising one or more (preferably two or more) (IIa) and (IIb), (a useful for producing a chimeric RNA produced comprises sequence to be transcomponents. The chimeric cell which involves concacting a target pre-mRNA expressed in the cell with (II) that is recognised by nuclear splicing components. The chimeric comprises nucleotide sequences comprising exons 1-10 of cystic preferably comprises nucleotide sequences comprising exons 1-10 of cystic fibrosis trans-membrane conductance regulator (GTRN). The chimeric RNA molecule produced using (II) which either comprises (A) or (B) comprises a nucleotide sequence to be trans-spliced to target pre-mRNA comprises and targeted cell death. (I) can be used for the treatment of comprises and targeted cell death. (I) can be used for the treatment of comprises and targeted cell death. (I) can be used for the treatment of comprises and diseases including genetic, infectious or autoimmune diseases and communistical produced ascener inventions or autoimmune diseases and communistical produced ascener inventions and to regulate gene expression in the present inventions the present interval.

Sequence 20 BP; 3 A; 6 C; 9 G; 2 T; 0 U; 0 Other;

exemplification of the present invention

Query Match 3.1%; Score 13; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 4.1e+02; Matches 13; Conservative 0; Mismatches 0; Indels

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Wed Apr 41 12:58:41 2004
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255 TCGGCCACGGTGC 267 ||||||||||||||||||1|| 15 TCGGCCACGGTGC 3

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ABL43850; ABL43850 RESULT

ABL43850 standard; DNA; 16 BP

11-APR-2002 (first entry)

Human chromosome 1p36-35 PCR primer SEQ ID NO:894.

chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome; PCR primer; ss. Human;

Homo sapiens.

JP2001321190-A.

20-NOV-2001.

12-MAR-2001; 2001JP-00068285

10-MAR-2000; 2000JP-00066716

(RIKA ) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.

WPI; 2002-144136/19.

Arraying genome clones.

Claim 4; Page 22; 528pp; Japanese.

The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the marker is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell cand lateral directions; (f) the mixed clones are cultured and the call three in the multiwell plates of the specified to midted respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified products; (h) the clones in the multiwell plates are specified from the amplified products; (h) the clones in the multiwell plates are specified from the amplified products; (h) the clones in the constituted as the positions on the chromosome and arrayed. The mixed reconstituted as the positions on the chromosome and arrayed. The mixed promess for human chromosome light-312 to ABL43531 to ABL43531 to PRE primers for human chromosome 21q22.1, which are specified as the present invention in the present invention 

Sequence 16 BP; 3 A; 7 C; 3 G; 3 T; 0 U; 0 Other;

Gaps . 0 Query Match

3.0%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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44 TGGCCACCACTCAGAG 59

8

1 TGGCCCCCACTCATAG 16

RESULT 437 AAQ47599/c ID AAQ47599 standard; cDNA to mRNA; 17 BP.

(revised)
(first entry) 25-MAR-2003 26-JAN-1994 AAQ47599;

7. 5117

Rat C RATRJG9/B-1258 jun-B specific probe.

Probe; quantification; human; GTP binding protein; G protein; alpha subunit; specific mENA; detection; hybridisation; diagnosis; pathophysiology; disease state; hereditary; cancer; infectious; osteodystrophy; pituitary tumour; acromegaly; melanoma cells; diabetes; PCR; polymerase chain reaction; ss.

Synthetic.

WO9315221-A1.

05-AUG-1993.

93WO-US000977 29-JAN-1993;

92US-00827208. 92US-00857059. 92US-00974409. 29-JAN-1992; 24-MAR-1992; 12-NOV-1992;

(HITB ) HITACHI CHEM CO LID. (HITB ) HITACHI CHEM RES CENT INC.

Akitaya T, Cooper A, Mitsuhashi M;

WPI; 1993-258695/32.

Quantitating messenger RNA in sample - using immobilised-polynucleotide having sequence complementary to sequence unique to the MRNA.

Example 9; Page 71; 177pp; English.

The sequences given in AAQ47594-603 show regions of homology between jun sequences and the jun-B specific probe B-1258 which may be of use as jun-B specific probes. They were used in the method of the invention for the getection and quantification of mRMAs in a sample without the need to detection and quantification of mRMAs in a sample without the need to purify the mRMA from cells. The claimed method comprises identifying a purify the mRMA from cells. The claimed method comprises identifying a polymucleotide sequence unique to the mRMA, and immobilising an oligomer components are washed from the support such that the unique sequence then incubated with the insoluble support such that the unique sequence components are washed from the support and bound RMA is labelled in such components are washed from the support. The amount of bound label is then a way that the label is incorporated onto the support relative to the amount of mRMA on the support. The amount of bound label is fine confortation of maltiple varieties of mRMA. It may be used for clearmined. This method can be used for the reliable, rapid, simultaneous conspanition of maltiple varieties of mRMA. It may be used for clearning and recognition of pathophysiology of various disease states, cancer, and infectious disease states. Contection is the molecular basis of hereditary of sprotein is the molecular basis of hereditary of contain mutant ds proteins. G proteins are also involved in invasive contain metastatic melanoma cells, and diabetes. See also AAQ473B1-666. 

Sequence 17 BP; 3 A; 8 C; 4 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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142 TGGCGGTGGAGGCCGG 157 

RESULT 438

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AAT50887-T50904 represent oligomucleotides of the invention. These sequences are all probes for interleukin-6 receptor (IL-6R) mRNA. IL-6 is one of the most well characterised of the cytokines. It functions through interacting with at least two transmembrane glycoprotein receptor molecules on the surface of target cells. The receptors are the IL-6R, and the signal transducer gp130. Signal transduction by IL-6 involves the concerted action of both IL-6R and gp130. IL-6 overproduction is implicated in many different disease states, particularly in cellular proliferation associated with these diseases. These sequences bind to the IL-6R coding sequence, thereby inhibiting IL-6R production. The sequences therefore inhibit the functioning of IL-6. These sequences can be used for inhibiting disease-associated cellular proliferation. The coligonucleotides are especially useful for treating cancer (e.g. renal cell carcinoma), autoimmune diseases or viral infections. They can also be used as probes for detecting IL-6 receptor mRNA, especially for evaluating the effectiveness of drugs in reducing IL-6 receptor mRNA
                                                                                                                                                    Probe, interleukin-6 receptor, IL-6R; cytokine, cellular proliferation, transmembrane glycoprotein receptor; signal transducer; gp130; inhibitor; IL-6; cancer; renal cell carcinoma; autoimmune disease; viral infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligo:nucleotide(s) complementary to interleukin-6 receptor mRNA - for treating proliferative diseases, e.g. cancer, auto-immune diseases or viral infections.

    .17
    /*tag= a
    /note= "optionally phosphorothioated"

                                                                                                                      Probe #3 for interleukin-6 receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Naidu YM;
                                                                                                                                                                                                                                                                             Location/Qualifiers
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AAT50889/c
ID AAT50889 standard; DNA; 17 BP.
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                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENP-) GEN-PROBE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-023093/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1995;
                                                                                                                                                                                                                                                                                                misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                      11-DEC-1996
                                                                                     26-AUG-1997
                                                                                                                                                                                                              therapy; ss
                                                                                                                                                                                                                                                                                                                                                                    EP747386-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brown SJ,
                                                                                                                                                                                                                                             Synthetic
                                                    AAT50889;
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Sequence 17 BP; 3 A; 7 C; 4 G; 3 T; 0 U; 0 Other;

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Gaps
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3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; cive 0; Mismatches 2; Indels
                              Conservative
                Local Similarity
                               14;
   Query Match
                Best Loca
Matches
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CGAGGGCCGCGCAGTG 89 N CGAGGACTCGCAGTG 74 17

ઠે 셤 RESULT 439 AAX68712/c

BP. AAX68712 standard; RNA; 17 (first entry) 28-JUL-1999 AAX68712;

Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like, tyrosine kinase 1; kinase insert domain containing receptor;

Human flt1 VBGF receptor hammerhead ribozyme substrate #7.

foetal liver kinase 1; ss.

WO9715662-A2.

Homo sapiens.

01-MAY-1997.

96WO-US017480, 25-OCT-1996; 95US-0005974P. 96US-00584040. 26-OCT-1995; 11-JAN-1996;

(RIBO-) RIBOZYME PHARM INC. (CHIR ) CHIRON CORP.

Escopedo J; Stinchcomb D, Pavco P, Mcswiggen J,

WPI; 1997-259017/23.

Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.

Claim 4; Page 46; 218pp; English.

The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGE). A patient (preferably human) having a condition associated with the level of the fms-like tyxosine kinase 1 (EL-1), kinase insert domain containing receptor (XDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour anglogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAXX7275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention 

Sequence 17 BP; 1 A; 6 C; 8 G; 0 T; 2 U; 0 Other;

ö 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; vative 0; Mismatches 2; Indels Matches 14; Conservative Query Match Best Local Similarity

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305 GAGCCCCGGGGACCGC 320 17 GAGCCCGGAGCCCGC 2 δ d

AAT85503 standard; cDNA; 17 AAT85503; RESULT 440 AAT85503
ID AAT8
XX AAT8
XX IT-N
XX IT

.. 0

BP.

(first entry) 17-NOV-1997 Oligo #13 used to isolate human chromosome 16 sequences.

Human; netrin; ATPase binding cassette transporter; ribosomal L3; augmenter of liver regeneration; hNET; hABC3; SEM L3; hALR; chromosome 16; exon trapping; axon; chicken; laminin domain; C. elegans;

GENZ ) GENZYME CORP

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The sequences given in AAT85503-06 are oligos which were used in the isolation of coding sequences from human chromosome 16. The invention contains details of the sequences encoding human netrin (NNET), human ATPASS Binding Osssette transporter (NABG3), human ribosomal L3 (SEM L3), and human augmenter of liver regeneration (NARR). The NNET gene can be used to develop chemostractants for use in axon regeneration. The hABG3 gene may be used in therapeutic applications for cystic fibrosis. The hABC3 gene may be used to develop products for treating damaged liver and liver diseases. The products can also be used for detection, diagnosis and screening assays. These oligonucleotides of may be used as primers in exon trap amplification experiments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, netrin; ATPase binding cassette transporter; ribosomal L3; augmenter of liver regeneration; NNET; hABC3; SEM L3; hALR; chromosome 16; exon trapping; axon; chicken; laminin domain; C. elegans; UNC-6; cystic fibrosis; ss.
                                                                                                                                                                                                                                                                                              New isolated human chromosome 16 genes - encode netrin, ATPase binding cassette transporter, ribosomal L3 sub-type or augmenter of liver
                                                                                                                                                                                                                      Connors TD, Dackowski WR, Klinger KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligo #1 hybridises to hABC3 cDNA sequence.
                                                                                                                                                                                                                                                                                                                                                              Disclosure, Page 18; 98pp; English
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UNC-6; cystic fibrosis; ss
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                                                                                                                                                                                                                      Burn IC,
                                                                                                                                                                                      (GENZ ) GENZYME CORP.
                                                                                                                                                                                                                                                                   WPI; 1997-108959/10
                                                                                                                                                                                                                                                                                                                                 regeneration.
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                                                                                                                           17-JUN-1996;
                                                                                                                                                          30-JUN-1995;
                                                                                                                                                                                                                      Landes GM, 1
Van Raay TJ;
                                                            WO9702346-A2
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                               Synthetic.
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                                                                                                                                                                                     The sequences given in AAPBS475-83 hybridise under stringent conditions to the sequence encoding the ATPase binding cassette transporter protein (hBABC3). The hABC3 genomic sequence was isolated from human chromosome 16 by exon trapping. hABC3 cDNA contains an open reading frame of 1685 amino acids. Comparison of ABC1, ABC2 and hABC3 reveals significant conservation in the regions surrounding the two ATP binding cassettes. The ATP binding cassettes of hABC3 flank a large linker domain containing numerous polar residues. The presence of these features in the linker domain suggests that this domain may play a regulatory role similar to the R domain of CPTR. The hABC3 gene may be used in therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; netrin; ATPase binding cassette transporter; ribosomal L3; augmenter of liver regeneration; hNET; hABC3; SEM L3; hALR; chromosome 16; exon trapping; axon; chicken; laminin domain; C. elegans; UNC-6; cystic fibrosis; ss.
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                                                                                      New isolated human chromosome 16 genes - encode netrin, ATPase binding cassette transporter, ribosomal L3 sub-type or augmenter of liver regeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                          Dackowski WR, Klinger KW;
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Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                             Connors TD,
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                                                                                                                                                                Claim 29; Page 61; 98pp; English.
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                             Burn TC,
                                                                         WPI; 1997-108959/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Landes GM, 1
Van Raay TJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JAN-1997.
                             Landes GM, |
Van Raay TJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT85480;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT85480
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**Page 213** 

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The sequences given in AAT85475-83 hybridise under stringent conditions to the sequence encoding the ATPase binding cassette transporter protein (hABC3). The hABC3 genomic sequence was isolated from human chromosome 16 by exon trapping. hABC3 cDNA contains an open reading frame of 1685 amino acids. Comparison of ABC1, ABC2 and hABC3 reveals significant conservation in the regions surrounding the two ATP binding cassettes. The ATP binding cassettes of hABC3 flank a large linker domain containing domain suggests that this domain may play a regulatory role similar to the R domain of CFTR. The hABC3 gene may be used in therapeutic

Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;

Gaps ö Query Match

3.0%; Score 12.8; DB 1; Length 17;

Best Local Similarity 87.5%; Pred. No. 3.2e+02;

Matches 14; Conservative 0; Mismatches 2; Indels

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RESULT 443 AAV95292/c ID AAV95292 standard; RNA; 17 BP.

AAV95292;

24-FEB-1999 (first entry)

Human c-fos target sequence nucleotide position 268.

Human; c-fos; hammerhead ribozyme; hairpin ribozyme; target site; cancer; oncogene; leukaemia; neuroblastoma; diagnosis; genetic drift; mutation; diseased cell; ss. 

Homo sapiens

WO9832846-A2.

30-JUL-1998

20-JAN-1998;

97US-0037658P. 97US-00998099. 23-JAN-1997; 24-DEC-1997;

(RIBO-) RIBOZYME PHARM INC

Jarvis I, Mcswiggen JA, Stinchcomb DT;

WPI; 1998-427942/36.

Enzymatic nucleic acid molecules which specifically cleave RNA derived from a c-fos gene - useful for treating conditions related to levels of c-fos, especially cancer.

Claim 2, Page 50, 72pp, English.

The present invention describes an enzymatic nucleic acid molecule which specifically cleaves RNA derived from a c-fos gene. AAV95401 to AAV95540 and AAV95541 to AAV95541 represent harmerhead ribozymes and hairpin ribozymes, respectively, which specifically cleave human c-fos. AAV95261 to AAV95400 and AAV95585 to AAV95628 represent human c-fos target cancer associated with elevated levels of c-fos oncogene, especially leukaemias, neuroblastomas and lung, breast and colon cancers. The ribozymes may also be used a diagnostic tools to examine genetic drift and mutations within diseased cells, or to detect the presence of c-fos RNA in a cell

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Gaps
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0
Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                         286 CCAAGCTGGTGAAGGA 301
                                                                                                                               17 ccardcredadadda 2
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Sequence 17 BP; 2 A; 7 C; 3 G; 0 T; 5 U; 0 Other;

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RESULT 444 AAV45792

AAV45792 standard; DNA; 17 BP.

AAV45792;

24-NOV-1998 (first entry)

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Primer NONA PCR-R.

Gene bank; combinatorial library; phagemid display; phage display; cosmixplexing; receptor; ligand; autoimmune disease; ss.

Synthetic.

WO9833901-A2

06-AUG-1998.

98WO-EP000533. 02-FEB-1998; 97EP-00101539 31-JAN-1997; (COSM-) COSMIX MOLECULAR BIOLOGICALS GMBH.

Collins J, Roettgen P;

WPI; 1998-437456/37.

Banks containing genes with restriction enzyme sites that generate specific cohesive ends - allowing production of large phage or phagemid display libraries, for screening to identify ligands for medical, diagnostic etc. use.

Example 1; Page 45; 87pp; English.

In a cosmiplexing method of the invention for the generation of double-stranded DNA inserts, the single-stranded hypervariable DNA oligos NONA-CA, NONA-CA, MONA-CA, 

Sequence 17 BP; 3 A; 4 C; 7 G; 3 T; 0 U; 0 Other;

Gaps Match 3.0%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 3.2e+02; les 14; Conservative 0; Mismatches 2; Indels of Query Match Best Local S

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262 CGGTGCACCTGGAGCA 277

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CGGGGTACCTGGAGCA 17

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Human chromosome 16 genes encoding netrin, ATP binding cassette transporter, ribosomal L3 and augmenter of liver regeneration proteins useful for, e.g. treatment of liver disease and cystic fibrosis.
                                                               Human, ATP binding cassette transporter, hABC3, cystic fibrosis,
treatment, trapping, modulation, expression, antibody, identification,
binding, substrate specificity, ligand, exon trap, PCR primer, amplify,
                                      Primer used to clone additional sequences from human ABC3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 40; Page 25; 220pp; English.
            03-JUN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                      (GENZ ) GENZYME CORP.
                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-063138/06.
                                                                                                                                                                                                                                                                          17-JUN-1996;
01-OCT-1996;
09-DEC-1996;
                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                16-JAN-1997;
                                                                                                                                                                                        WO9748797-A1
                                                                                                                                                                                                                  24-DEC-1997.
                                                                                                                                                                                                                                                                                                                                                                  Landes GM,
Klinger KW;
                                                                                                                                            Synthetic.
<u>(</u>,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotides AAV16310-25 are used to clone addional sequences of nucleic acids encoding human netrin (hNET), human ATP binding cassette transporter (hAEC3), human ribosomal 13 (RD121), and human augmenter of liver regeneration (hALR). Partial DNA sequences from these genes were isolated using exon traps AAW46753-57. Genetrapper, 3' RACE and RT-PCR were employed to identify additional sequences. The antisense coligonuclectides of the isolated sequences are used to modulate expression of hNET, hABC3, RP131 or hALR, and prevent its translation. Antibodies against hNET, hABC3, RP131 and hALR can be used to block binding of their naturally occurring ligands. The host cells containing vectors with DNA inserts encoding the proteins can be used to a method for identifying compounds which bind to hNET, hABC3, RP131 or hALR. Modulation or alteration of hABC3 substrate specificity may have significant therapeutic implications for cystic fibrosis. hALR could be used in the treatment of damaged liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human chromosome 16 genes encoding netrin, ATP binding cassette transporter, ribosomal L3 and augmenter of liver regeneration proteins useful for, e.g. treatment of liver disease and cystic fibrosis.
                                                                                                                                            Human; netrin; hNET; ATP binding cassette transporter; hABC3; ribosomal L3; RPL3L; augmenter of liver regeneration; hALR; treatment; trapping; modulation; expression; antibody; identification; binding; substrate specificity; ligand; exon trap; PCR primer; amplify; ss.
                                                                                                                Primer used to clone additional sequences of ABC, NET, ALR and RPL3L.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Connors TD, Dackowski WR, Van Raay TJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ch 3.0%; Score 12.8; DB 1; Length 17; 1 Similarity 87.5%; Pred. No. 3.2e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 35; Page 24; 220pp; English.
                                                                                                                                                                                                                                                                                                                                                       96US-00665259.
96US-00720614.
96US-00762500.
                                AAV16316 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                            97WO-US000785
                                                                                        03-JUN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Burn TC,
                                                                                                                                                                                                                                                                                                                                                                                                                (GENZ ) GENZYME CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-063138/06.
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                       17-JUN-1996;
01-OCT-1996;
09-DEC-1996;
                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                    WO9748797-A1.
                                                                                                                                                                                                                                                                                                                            16-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Landes GM,
Klinger KW;
                                                                                                                                                                                                                                                                                                24-DEC-1997
                                                                                                                                                                                                                       Synthetic.
                                                            AAV16316;
    RESULT 445
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Dackowski WR, Van Raay TJ;

Burn TC, Connors TD,

96US-00665259. 96US-00720614. 96US-00762500.

97WO-US000785.

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oligonuclectides AAV16326-32 are used to clone addional sequences from nucleic acids encoding human ATP binding cassette transporter (hABC3).

Partial DNA sequences from the genes was isolated using exon traps additional bequences. The ABC gene is located in the PKD1 locus, between additional sequences. The ABC gene is located in the PKD1 locus, between the sequence shows homology with murine ABC1 and ABC2 genes. The ABC proteins are responsible for the transport of a wide variety of substrates across cell membranes. Proteins and across cell membranes. Proteins and across subcellular strong structural similarities. ABC transporters govern unidirectional cransport of molecules into or out of cells and across subcellular membranes. The antisense oligonucleotides of the ABC3 gene sequence are membranes. The antisense oligonucleotides of the ABC3 gene sequence are commented to modulate expression of ABC prevent its translation. Antibodies against ABC can be used to block binding of its naturally occurring cagains the cells containing vectors with DNA inserts encoding the C protein can be used in a method for identifying compounds which bind to the ABC. Modulation or alteration of hABC3 substrate specificity may have be significant therapeutic implications for cystic fibrosis
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Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 4 A; 3 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          288 AAGCTGGTGAAGGACC 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA36411/c

ID AAA36411 standard; DNA; 17

XX

AC AAA36411;

XX

DT 26-JUL-2000 (first entry)
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Gaps

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288 AAGCTGGTGAAGGACC 303

AAV16329 standard; DNA; 17

RESULT 446

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AAV16329;

AAV16329 ID AAV1 XX AC AAV1

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Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis; allele specific ollogonucleotide; ASO; reduced complexity genome; RCG; genomic classification; identification; DNA fingerprinting; tumour characterisation; hybridisation; 88.
                                                                                                                                                                                                                                                               Detection of single nucleotide polymorphisms in genomes by prepara and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs.
         Human genomic SNP allele specific oligonucleotide SEQ ID NO:477
                                                                                                                                                                                                                      Charest A;
                                                                                                                                                                                                 (MASI ) MASSACHUSETTS INST TECHNOLOGY
                                                                                                                                                                                                                      Housman DE,
                                                                                                                                                                                                                                                                                                              Disclosure; Page 67; 111pp; English.
                                                                                                                                                      99WO-US022283
                                                                                                                                                                          98US-0101757P
                                                                                                                                                                                                                        Landers JE, Jordan B,
                                                                                                                                                                                                                                             WPI; 2000-293181/25.
                                                                                                          WO200018960-A2.
                                                                                      Homo sapiens
                                                                                                                                                      24-SEP-1999;
                                                                                                                                                                            25-SEP-1998;
                                                                                                                                 06-APR-2000
```

A method has been developed for detecting the presence or absence of a single nucleotide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RCG) from the genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent sequences used in the exemplification of the present invention. AAA35948 to

Sequence 17 BP; 1 A; 7 C; 2 G; 7 T; 0 U; 0 Other;

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3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                     Local Similarity 87.5
es 14; Conservative
       Query Match
                         Best Loca
Matches
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AAF02688 standard; DNA; 17 BP AAF02688; RESULT 448 AAF02688/C 

(first entry) 16-FEB-2001

Hammerhead ribozyme substrate #983.

Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.

Homo sapiens

WO200061729-A2

19-OCT-2000

11-APR-2000; 2000WO-US009721

12-APR-1999;

(RIBO-) RIBOZYME PHARM INC.

Mcswiggen J; Zwick M, Pavco P, Blatt L, 

WPI; 2000-647423/62.

Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.

Claim 37; Page 78; 164pp; English.

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor. EAR3/COUP-TP-1, the GATA transcription factor gene, IRF-2 and/or the GAAT Displacement Protein (CPP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and interferon alpha

Sequence 17 BP; 0 A; 9 C; 4 G; 4 T; 0 U; 0 Other;

preparation

ò Gaps ; Length 17; 2; Indels ).0%; Score 12.8; DB 1; Local Similarity 87.5%; Pred. No. 3.2e+02; les 14; Conservative 0; Mismatches 2; Query Match Best Loca Matches

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AAF05332 standard; DNA; 17 BP. RESULT 449 

AAF05332;

(first entry) 16-FEB-2001

Ribozyme, erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss. Hammerhead ribozyme substrate #2551.

Homo sapiens

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WO200061729-A2.

19-OCT-2000.

11-APR-2000; 2000WO-US009721.

99US-0129390P 12-APR-1999;

Blatt L, Zwick M, Pavco P, (RIBO-) RIBOZYME PHARM INC.

Mcswiggen J;

WPI; 2000-647423/62.

Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.

Claim 18; Page 114; 164pp; English

The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes

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99US-0129390P

Oligonucleotide array; genotyping; single base extension reaction; PCR primer; polymorphic locus; single nucleotide polymorphism; ss.

Oligonucleotide

Reverse primer #67 used in multiplexing PCR/SBE assay

(first entry)

02-FEB-2001

AAC73338;

AAC73338 standard; DNA; 17 BP.

RESULT 4

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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRE-2 and/or the CAATI Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoletin, granulocyte colony stimulating factor protein and interferon alpha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ensymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
   the GATA transcription
encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production erythropoietin, granulocyte colony stimulating factor protein and interferon alpha
                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozyme; erythropoietin; granulocyte colony stimulating factor; interferon alpha; ss.
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3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                         Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                        Sequence 17 BP; 1 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Blatt L, Zwick M, Pavco P, Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hammerhead ribozyme substrate #1181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 37; Page 82; 164pp; English.
                                                                                                                                                                                                                                                                                                                                              886/c
AAF02886 standard; DNA; 17 BP.
                                                                                                                                                                                                                                  345 CGGCTGCTCTACAGCG 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-APR-2000; 2000WO-US009721.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0129390P.
                                                                                                                                                                                                                                                             GGCCTGCTCTTCAGCG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                 16-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                             AAF02886;
                                                                                                                                                                                                                                                                                                                         RESULT 450
AAF02886/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNF). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to
                                                                                                                                                                                                                                                                                                                                                                                                                             Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                          5
                                                                                                                                                                                                                                                                                                                                                            Lockhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                            Kaplan P, Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                  (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 7; Page 55; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                             Hirschhorn JN, Huang X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     392 CGCCAAGAAGGTCTTC 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CGCCAACATGGTCTTC 17
                                                                                                                                                                                                                                                                         99US-0126473P.
99US-0140359P.
                                                                                                                                                                                                                                           27-MAR-2000; 2000WO-US008069.
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Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                                                                                             Fan J, Hirschhorn
Ryder T, Sklar P;
                                                                                                                                                                                     WO200058516-A2.
                                                                                                                                                                                                                                                                         26-MAR-1999;
23-JUN-1999;
                                                                                                                                                          Unidentified.
                                                                                                                                                                                                                 05-OCT-2000.
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Gaps

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GCACCTGGAGCAGGGC 281

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GCACCGGGAGCGGGGC 1

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DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; martle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens. Synthetic.

WO200159103-A2.

16-AUG-2001

09-FEB-2001; 2001WO-US004273

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBOZYME PHARM INC BLATT L. (RIBO-) H (BLAT/) H (MCSW/) N

Chowrira BM Blatt L, Mcswiggen J, (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 79; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NG20). The crequlates expression of a neurite growth inhibitor gene (NG20). The collar acids may be enzymatic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NRNA motif) propersessing an NCH motif) a G-cleaver (cleaving RNA with a NRNA motif) propersessing an NCH motif), a G-cleaver (cleaving RNA with a NCH motif) propersessing an NCH motif), a G-cleaver (cleaving RNA with a NCH motif) propersessing an NCH motif) a g-cleaver (cleaving RNA with a SPA) a sinzyme (cleaving RNA with a SPA) and the presence of a divalent cation that is preferably MG<sup>2</sup>+.

Furthermore, it may be contacted with a coll to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of fone or more chargeting nucleic acid may be contacted with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably MG<sup>2</sup>+. Furthermore, the condition associated MEL, mantle-cell and treat a patient having a condition associated Mel, the level of collicular space or follicular space and the cell and treat a patient having a condition associated with the level of the condition associated with the level of the sapeies. In particular, the NOGO-targeting nucleic acid may be used to treat comprise the use of one or more content and patient having a condition associated with the level of treat central nervous system (NNS) injury model cation and central nervous system (NNS) injury model acceptovascular accident for ender a patient having a condition associated with the level of the arbity amyotrophic lateral acceptorate and advanced a disease, ataxia, Huntington's disease, central response in the model s sequence is an inozyme of the invention

Sequence 17 BP; 1 A; 8 C; 8 G; 0 T; 0 U; 0 Other;

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                                                                                                                                                                                                                          Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; declarer; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; lymphocytic leukaemia; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
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0
Length 17;
3.0%; Score 12.8; DB 1; Length 1
87.5%; Pred. No. 3.2e+02;
cive 0; Mismatches 2; Indels
                                               302 CCTGAGCCCCGGGGAC 317
                                                                  2 CCGGCCCCCGGGGAC 17
                                                                                                                                ABK02394 standard; RNA; 17
                                                                                                                                                                                12-MAR-2002 (first entry)
3.0%
Best Local Similarity 87.5%
Matches 14; Conservative
                                                                                                                                                                                                       Human NOGO Amberzyme #66.
                                                                                                                                                         ABK02394;
                                                                                                         RESULT 453
                                                                                                                       ABK02394
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WO200159103-A2. 16-AUG-2001.

Homo sapiens.

Synthetic.

09-FEB-2001; 2001WO-US004273

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Blatt L, Mcswiggen J, Chowrira BM;

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 131; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NOGO). The nucleic acids (e.g. a ribozyme or a DNAZYME) and nozyme (an endolytic nucleic acids (e.g. a ribozyme or a DNAZYME) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaving RNA with a NYM motif) pr an amberzyme (cleaving RNA with a NGM triplet), a zinzyme (cleaving RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2^+. Purthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more

treat lymphoma, leukaemia, be-cell lymphoma, low-grade or follicular non-thedgkin's lymphoma (NLL), bulky low-grade or follicular NLL, lymphocytic leukaemia, leukaemia, be-cell lymphoma, low-grade or follicular NLL, lymphocytic leukaemia, HIV (human immunodeficiency virus) associated NLL, mantle-cell lymphoma (NCL), immunocytoma (INC), small b-cell lymphocytic lymphoma, (NCL), immunocytoma (INC), small b-cell lymphocytic lymphoma, compared to cleave RNA of the NoGO gene in the presence of a divalent cation that is preferably MG-7-. Furthermore, the nucleic acid may be contacted with a cell to reduce NoGO activity of the nucleic acid may be contacted with a cell to reduce NoGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CNA, stroke), Alzahmar's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MLS), parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, dasses, muscasse, muscasse, muscasse, creates which respond to the modulation of NOGO expression. The present sequence is an amberzyme molecule of the invention

Sequence 17 BP; 1 A; 9 C; 7 G; 0 T; 0 U; 0 Other;

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Gaps
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0
ch 3.0%; Score 12.8; DB 1; Length 17; I Similarity 87.5%; Pred. No. 3.2e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                     305 GAGCCCCGGGGACCGC 320
                                                                                                   GCGCCCCGGGGACCCC 17
                   Local Similarity
       Query Match
                       Best Loca
Matches
                                                                         ò
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ABK01169 standard; RNA; 17 BP 12-MAR-2002 (first entry) Human NOGO Inozyme #439. ABK01169; RESULT 454 ABK01169, 

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; noctropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; g-cleaver; amberzyme; inizyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebrovascular accident; cVA; Alzheimer's disease; multiple sclerosis; Parkinson's disease; ataxia; Huntingcon's disease; ceretzeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

40200159103-A2. sapiens 6-AUG-2001. Synthetic Ношо

BLATT L. MCSWIGGEN J. CHOWRIRA B M. CHOM/)

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. 39-FEB-2001; 2001WO-US004273. RIBOZYME PHARM INC. RIBO-) (BLAT/) Chowrira BM; Mcswiggen J, Blatt L,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 85; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGOD). The regulates expression of a neurite growth inhibitor gene (NGOD). The nucleic acids (e.g. a ribozyme or a nucleic and in nozyme (an endolytic nucleic acid cleaving RN with a NTM motif) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NCH will) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NCH will). The CD20-targetting nucleic acid is used to cleave RNA of the presence of a divalent cation that is preferably MG<sup>2</sup> +. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular nontraced with a cell low-grade or follicular nontraced with a cell lymphocytic lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, conductory arthropathy. The NOGO-tergetting nucleic acid may be contacted with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably MG<sup>2</sup> +. Furthermore, the condition associated with the level of the treatment may further comprise the use of one or more cell and treat a patient having a condition associated with the level of the stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemchbrapy-induced neuropathy, and/cortexpic lackal acid may be used to treat central nervous system (CNG) injury and cerebrovascular accident CCC theratosh's disease, ataxia, Huntington's disease, cataxia, Huntington's disease, cataxia, Huntington's disease, cataxia, Huntington's disease, mescular disease, cataxia, Huntington's disease, ataxia, Hunting sequence is an inozyme of the invention

Sequence 17 BP; 2 A; 6 C; 2 G; 0 T; 7 U; 0 Other;

Gaps ö 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; vative 0; Mismatches 2; Indels Query Match Best Local Similarity 87.5 Matches 14; Conservative

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ABK00842 standard; RNA; 17 BP. 286 CCAAGCTGGTGAAGGA 301 (first entry) Human NOGO Inozyme #112. 12-MAR-2002 ABK00842; RESULT 455
ABK00842
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AC ABK0084
DT 12-MARXX
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HUMAN 10
KW MUSCUL
KW MUS g ઠ

Human; 88; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; deleaver; amberzyme; zinzyme; lymphoma; leukaemia; human immunodeficiency virus; HV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immunocytoma; IMC; immunocytoma; IMC; immunocytoma; MC; immunocytoma; IMC; immunocytoma; imc; immunocytoma; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS;

ABK02395 standard; RNA; 17 BP.

ABK02395;

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (MGO). The regulates expression of a neurite growth inhibitor gene (MGO). The concleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids an NCH montif), a G-cleaver (cleaving RNA with an NCH motif), a g-cleaver (cleaving RNA with a NCH motif), a zinzyme (cleaving RNA with a Presence of a divalent cation that is preferably MG<sup>2</sup>+.

CC of CD20 in the presence of a divalent cation that is preferably MG<sup>2</sup>+.

Creat Lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular nonce (c CD20). The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma (MCL), immunocytoma (NH), bulky low-grade or follicular NH). Immunocytoma (NH), bulky low-grade or follicular noncentageting nucleic acid may be contacted with a cell to reduce NOGO gene in the presence of a divalent cation that is preferably MG<sup>2</sup>+. Furthermore, the creatment may divident acid may be contacted with a cell to reduce NOGO activity of the contacted may be contacted with a cell to reduce NOGO activity of the contacted may be contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the cell and treat a patient having a contacted with a cell to reduce NOGO activity of the contacted w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and
Parkinson's disease, ataxia, Huntington's disease,
Creutzfeldt-Jakob disease, muscular dystrophy, neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence is an inozyme of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chowrira BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 88; Page 79; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        central nervous system injury.
                                                                                                                                                                                                                                                                              11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                                                                                                                                  09-FEB-2001; 2001WO-US004273
                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                         (CHOW/) CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                    BLATT L.
MCSWIGGEN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-607195/69.
                                                                                                                                       WO200159103-A2.
                                                                       sapiens
                                                                                                                                                                                     16-AUG-2001
                                                                                               Synthetic.
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Sequence 17 BP; 1 A; 9 C; 7 G; 0 T; 0 U; 0 Other;

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids aribozyme or a conclain and beaving an expression of an entrangent of cleaving an expression and motify in a G-cleaver (cleaving RNA with a NYN motif) proposessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) and a makezyme (cleaving RNA tith an NYN triplet), a zinzyme (cleaving RNA with a NYN motif). The CD20-targetting nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA coff could be contacted with a cell to reduce CD20 eactivity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more chargetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-cleaked min and inflammatory virus) associated NHL, lymphocytic lymphoma, lymphoma, lymphoma, limin minimus (NHL), mantle-cell lymphoma (NCL), immunocytoma (NMC), small B-cell lymphocytic lymphoma, immune thrombocytopeania, and inflammatory arthropathy. The NOGO gene in the

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                                                 Gaps
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0
Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                   14; Conservative
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303 CTGAGCCCCGGGGACC 318 

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carebroprotective, notropic; ortobrates, antipamimatory; inamication; carebroprotective; notropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; NDXzyme; informa; linozyme; informa; linozyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin; lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; huC1; immunocytoma; MC1; immune thromborytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; Parkinson's disease; ataxia; Huntington's disease; creuzieldt-Jakob disease; muscular dystrophy; neurodegenerative disease. Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury. therapy; cytostatic; antiinflammatory; haemostatic; Chowrira BM; Claim 88; Page 131; 200pp; English. 11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P. 09-FEB-2001; 2001WO-US004273. RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (first entry) Human NOGO Amberzyme #67. Blatt L, Mcswiggen J, CHOWRIRA B M. Human; ss; antisense WPI; 2001-607195/69. BLATT L. MCSWIGGEN J. WO200159103-A2. sapiens. 12-MAR-2002 16-AUG-2001. Synthetic. CHOM/) (MCSM/) Ношо 

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886666666666688888
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presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more treat central nervous system (CMS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), Parkinson's disease, ataxia, Huntington's disease, Creuzzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present sequence is an amberzyme molecule of the invention

Sequence 17 BP; 1 A; 9 C; 6 G; 0 T; 1 U; 0 Other;

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2; Indels 0; Gaps
Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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305 GAGCCCCGGGGACCGC 320 1 GCGCCCGGGGACCCC 16 ò д

457 RESULT

ABN07567 standard; DNA; 17 BP ABN07567; ABN07567 

29-MAY-2002 (first entry)

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7559.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
21-SEP-2000; 2000US-0234559P.
04-0CT-2000; 2000US-0024559P.
04-0CT-2000; 2000US-002651.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.

(AEOM-) AEOMICA INC.

Shannon ME Chen W, Rank DR, Hanzel DK, Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

2001WO-US000668. 2001WO-US000669. 2001WO-US000670. 2001US-0266860P.

30-JAN-2001; 30-JAN-2001; 05-FEB-2001;

Disclosure; SEQ ID NO 7559; 214pp; English

The present invention describes a human genome-derived myosin-like protein 1 (hgbMLP-1). The protein and polymucleotide sequences of hgbMLP-1 can be used in gene therapy and vaccine production. The hgbMLP-1 nucleic acids in samples, as amplification substrates, to nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hgbMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hgbMLP-1 proteins or polypeptides may be expressing the proteins. The hgbMLP-1 proteins or polypeptides may be concentrated in a standards in assays used to determine the concentration and/or amount specifically of hgbMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desorption indisation, as the present in patients having specific deficiency in hgbMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration and skeletal muscle disorders. hgbMLP-1 may be used for diagnosing a classification while the expression of hgbMLP-1, in particular heart of hgbMLP-1 sequence in the exemplification of the present invention. W.B. The sequence data for this patent did not form part of the printed sequence are this patent did not form part of the printed at the present invention. The sequence of the sequence of the sequence of the sequence of the present invention. W.B. The sequence data for this patent did not form part of the printed sequence. ô Human, genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1, heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; 88. Gaps Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5992. ö Query Match
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 6 A; 3 C; 6 G; 2 T; 0 U; 0 Other; 26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234667P.
27-SEP-2000; 2000US-0235359P.
04-0CT-2000; 2000US-0024563.
30-JAN-2001; 2001W0-US000662.
30-JAN-2001; 2001W0-US000663.
30-JAN-2001; 2001W0-US0006663.
30-JAN-2001; 2001W0-US0006663.
30-JAN-2001; 2001W0-US0006665. ABN06000 standard; DNA; 17 BP. 385 ACGACGGCGCCAAGAA 400 25-MAY-2001; 2001WO-US016981 2 ATGACGGGCCAAGAA 17 29-MAY-2002 (first entry) WO200192524-A2. Homo sapiens. 06-DEC-2001. ABN06000; RESULT 458 ABN06000/ X56666666666666666666666666888 ઠ 셤

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The present invention describes a human genome-derived myosin-like

protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1

c can be used in gene therapy and vaccine production. The hGDMLP-1

nucleic acids can be used as probes to detect, characterise and quantify

c hGDMLP-1 nucleic acids in samples, as amplification substrates, to

provide initial substrates for the recombinant engineering of hGDMLP-1

protein variants having desired phenotypic improvements, and for

expressing the proteins. The hGDMLP-1 proteins or polypeptides may be

crapture proteins, as standards in assays used to determine the concentration

and/or amount specifically of hGDMLP proteins, as specific biomolecule

capture probes for surface-enhanced laser description ionisation, as

therapeutic supplement in patients having specific deficiency in hGDMLP-1

production, and in vaccines or for replacement therapy. The

colynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a

colynucleotide sequence represents an oligomer used in the screening of the

condisorders associated with the expression of hGDMLP-1. in particular heart

cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence in the expression of the present invention. N.B.

Crapture probes for this patent did not form part of the printed

specification, but was obtained in electronic format directly from WIPO

cat fitp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                        New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
Shannon ME;
Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 5 A; 2 C; 7 G; 3 T; 0 U; 0 Other;
     Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 5992; 214pp; English
     Ji Y, Penn SG,
                                                                                                            WPI; 2002-179446/23.
     Gu Y,
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Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                  351 CTCTACAGCGACTICC 366
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Gaps

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Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                        Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:5988.
ABN05996 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                            26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-023468PP.
27-SEP-2000; 2000US-023559P.
04-0CT-2000; 2000GB-00024263.
30-JAN-2001; 2001WO-US000661.
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                                                   29-MAY-2002 (first entry)
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                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                            06-DEC-2001.
                         ABN05996;
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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 can be used as probes to detect, characterise and quantify concleic acids can be used as probes to detect, characterise and quantify concleic acids can be used as probes to detect, characterise and quantify concleic acids the samples, as amplification substrates, to protein variants having desired phenotypic improvements and for capturessing the proteins. The hGDMLP-1 proteins or polypeptides may be concentration to and/or amount specifically of hGDMLP-1 proteins, as specifically recognise hGDMLP-1 proteins, as specifically of hGDMLP-1 proteins, as specific blomolecule and/or amount specifically of hGDMLP-1 proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as capture probes for surface-enhanced laser desorption ionisation, as production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The conclust associated with the expression of hGDMLP-1, in particular heart of isorders associated with the expression of hGDMLP-1, in particular heart can skeleral muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed capture of the present invention. N.B. concentration, but was obtained in electronic format directly from MIPO capture.
                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                   Shannon ME;
                                                                                                                                                                                                                                                                                                                      Chen W,
                                                                                                                                                                                                                                                                                                                   Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure, SEQ ID NO 5988, 214pp, English.
30-JAN-2001; 2001WO-US000662.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
                                                                                                                                                                                                                                                                                                                      Gu Y, Ji Y, Penn SG,
                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                       (AEOM-) AEOMICA INC.
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0; Gaps Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels 354 TACAGCGACTTCCTCA 369 ઠ

Sequence 17 BP; 6 A; 2 C; 5 G; 4 T; 0 U; 0 Other;

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BP 17 TACATGGACTICCTCA 2 ABN01017 standard; DNA; 17 RESULT 460 ABN01017 셤

ABN01017;

Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1009. (first entry) 29-MAY-2002 X S X E X E X E X S X S X S X S

Human, genome-derived myosin-like protein 1, GDMLP-1; hGDMLP-1; heart, muscle, myosin, chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 claim be used as probes to detect, characteries and duantify a nucleic acids an beamples, as amplification substrates for the recombinant engineering of hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypoptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The production and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fip.wipo.int/pub/published_pot_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen W, Shannon ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 8 A; 3 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 1009; 214pp; English.
                                                                                                                                                                                                                                                                                                                              30-JAN-2001, 2001WO-US00661, 30-JAN-2001, 2001WO-US006661, 2001WO-US006661, 2001WO-US000661, 2001WO-US00061, 2001WO-US0006
                                                                                                                                                                                                                                    2000US-0234687P,
2000US-0236359P,
2000GB-00024263,
                                                                                                                                   25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-179446/23.
WO200192524-A2.
                                                               06-DEC-2001
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Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

202 CGGTGAAAGCAGAA 217

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ABN01018 standard; DNA; 17 BP. ABN01018; RESULT 461 ABNO1018 ID ABNO1001 XX AC ABNO1001

Human, genome-derived myosin-like protein 1, GDMLP-1, hGDMLP-1, heart, muscle, myosin, chromosome 22, gene therapy, vaccine, heart disease, skeletal muscle disorder, amplicon, screening; ss. Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1010. 27-5EP-2000; 200045-023489; 27-5EP-2000; 200045-023489; 30-4AN-2001; 2001MG-US000661. 30-4AN-2001; 2001MG-US000663. 30-4AN-2001; 2001MG-US000664. 30-4AN-2001; 2001MG-US000666. 30-4AN-2001; 2001MG-US000666. 30-4AN-2001; 2001MG-US000669. 30-4AN-2001; 2001MG-US000669. 30-4AN-2001; 2001MG-US000669. 30-4AN-2001; 2001MG-US000669. 30-4AN-2001; 2001MG-US000669. 25-MAY-2001; 2001WO-US016981 05-FEB-2001; 2001US-0266860P (first entry) 40200192524-A2. Homo sapiens. 26-MAY-2000; 29-MAY-2002

Shannon ME; Gu Y, Ji Y, Penn SG, Hanzel DK, Rank DR, Chen W, WPI; 2002-179446/23.

(AEOM-) AEOMICA INC.

New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure; SEQ ID NO 1010; 214pp; English.

The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify nucleic acids can be used as probes to detect, characterise and quantify herbit and lead to the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specifically recognise hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser describing as specific deficiency in hGDMLP-1 proteints, as specific deficiency in hGDMLP-1 proteints, as specific deficiency in hGDMLP-1 proteints associated with the expression of hGDMLP-1, in particular heart of sequence in the expression of hGDMLP-1, in particular heart and in was disorders. hGDMLP-1 is localised to chromsome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence date for this partent did not form part of the printed sequence date for this patent did not form part of the printed sequence cate for this patent did not form part of the printed sequence at five with the leavence of the present sequence at the leavence of the present sequence at the leavence of the present invention. N.B.

The sequence date for this patent did not form part of the printed sequence at the land of the present did not form part of the printed by the was obtained in electronic format directly from WIPO at the present sequence.

Sequence 17 BP; 8 A; 2 C; 7 G; 0 T; 0 U; 0 Other;

Gaps ô Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels

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Wed Apr 41 14:56:41 400%
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CAGGGAAAGCAGAGAA 16 202 CGGTGAAAGCAGAGAA ч

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ABN07571 standard; DNA; 17 29-MAY-2002 (first entry) ABN07571; RESULT 4 

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:7563.

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234587P.
04-OCT-2000; 2000US-02036359P.
04-OCT-2000; 200UMO-US000661.
30-JAN-2001; 200UMO-US000662.
30-JAN-2001; 200UMO-US000663.
30-JAN-2001; 200UMO-US000666.
30-JAN-2001; 200UMO-US000666.
30-JAN-2001; 200UMO-US000666.
30-JAN-2001; 200UMO-US000666.
30-JAN-2001; 200UMO-US000666.

(AEOM-) AEOMICA INC.

DK, Rank DR, Chen W, Shannon ME; Hanzel Gu Y, Ji Y, Penn SG, WPI; 2002-179446/23. New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser descrption ionization, comprises human myosin-like protein hGDMLP-1.

Disclosure, SEQ ID NO 7563; 214pp; English.

The present invention describes a human genome-derived myosin-like

protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP
c l can be used in gene therapy and vaccine production. The hGDMLP-1

c nucleic acids can be used as probes to detect, characterise and quantify

nucleic acids can be used as probes to detect, characterise and quantify

c provide initial substrates for the recombinant engineering of hCDMLP-1

c protein variants having desired phenotypic improvements, and for

expressing the proteins. The hGDMLP-1 proteins or polypeptides may be

cused as immunogens to raise antibodies that specifically recognise hGDMLP
c used as immunogens to raise antibodies that specifically recognise hGDMLP
c and/or amount specifically of hGDMLP-proteins, as specific biomolecule

c and/or amount specifically of hGDMLP proteins, as specific biomolecule

c capture probes for surface-enhanced laser desorption indisation, as

therapeutic supplement in patients having specific deficiency in hGDMLP-1

c production, and in vaccines or for replacement therapy. The

polymuloeride sequences encoding hGDMLP-1 may be used for diagnosing a

c disorder associated with the expression of hGDMLP-1, in particular heart

and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hobWip-1 sequence in the exemplification of the present invantion. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequence ន្តន្តន្តន្តន្ត

Sequence 17 BP; 6 A; 3 C; 7 G; 1 T; 0 U; 0 Other;

.; 0 Gaps .. 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; rative 0; Mismatches 2; Indels Best Local Similarity 87.5 Matches 14; Conservative Query Match

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ABK26660 standard; DNA; 17 BP.

ABK26660;

09-APR-2002 (first entry)

Waxy starch production genome altering oligonucleotide #316.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycinh. Brimariabiotic stress tolerance; impromental tolerance; hygromycinh. Brimariand acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; way starch; altered floral morphology; male-sterile plant; albino mutant; and fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced innolenic acid production; RESULT 46

ABK26666

XX

XX

ABK26666

DT

O9-A

CABC2

XX

ABK2

ABK2

ABK2

ABK3

Oryza sativa. Synthetic. WO200192512-A2.

06-DEC-2001.

01-JUN-2001; 2001WO-US017672.

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

(UYDE ) UNIV DELAWARE.

Rice MC, Kim Kmiec EB, Gamper HB,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 163; 220pp; English.

The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications consist of o-methyl modification, an LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or

directing repair or alterations. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The coligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abbotic stress tolerance, improved contritional value (e.g. altering amino acid content of plants or conferring amino acid over production, herbicide resistance (e.g. offersting amino acid over production), herbicide resistance (e.g. grasistance, porphyric herbicide resistance or triazine resistance), disease resistance, nondified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral corphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid). The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome callering oligonucleotide of the invention

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Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

Gaps . 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; ative 0; Mismatches 2; Indels Local Similarity 87.5 nes 14, Conservative Query Match Matches

380 CCGCGACGACGCCCC 395 2 cadddadradddddd 17 ठे g

5639/c ABK26639 standard; DNA; 17 BP. 09-APR-2002 (first entry) ABK26639; ABK26639, 

Waxy starch production genome altering oligonucleotide #295.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin; abiotic stress tolerance; improved mutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; imidazollinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; waxy starch; altered floral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced plant; albino mutant; increased stearate production; reduced linolenic acid production; abino plant; increased stearate production; reduced linolenic acid production; photosynthetic process.

Oryza glaberrima.

WO200192512-A2.

06-DEC-2001

01-JUN-2001; 2001WO-US017672

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

(UYDE ) UNIV DELAWARE.

Kmiec EB, Gamper HB, Rice MC,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 162; 220pp; English.

The invention relates to an oligomuclectide for targeted alteration of a genetic sequence, which comprises a single-stranded oligomuclectide featuble a DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligomuclectide. The chemical modifications of the oligomuclectide. The chemical modification is LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or more of these modifications. The oligomuclectides are useful for configuration repair or alteration of plant genetic information. The oligomuclectides are particularly useful for creating plants with desired opencypes, e.g. environmental or abiotic stress tolerance, improved conferring amino acid over production), herbicide resistance, improved conferring amino acid over production), herbicide resistance, conferring amino acid over production), herbicide resistance, conferration of plants of plants of glyphosate resistance, indified oil production, modified fatry acid content (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatry acid content of e.g. increased starch or production of waxy starch), altered floral corpusional plants of photosynthetic processes. This sequence represents a genome analysis of photosynthetic processes. This sequence represents a genome content of plants of the invention 

Sequence 17 BP; 1 A; 6 C; 7 G; 3 T; 0 U; 0 Other;

Gaps .; 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels Query Match Best Local Similarity 87.5' Matches 14; Conservative

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ABK26659 standard; DNA; 17 BP.

09-APR-2002 (first entry) ABK26659;

Waxy starch production genome altering oligonucleotide #315.

Chromosomal genomic alteration; genome altering oligonucleotide, PCR; ss, o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA atteration; environmental tolerance; hydromychin-b; abiotic stress tolerance; improved nutritional value, hydromychin; primer; amino acid over production; herbicide resistance; glyphosate resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistence; disease resistance; altered floral morphology; male-sterile plant; albino mutant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; increased stearate production; reduced alinolenic acid production; photosynthetic process. 

Oryza sativa. Synthetic.

WO200192512-A2.

06-DEC-2001.

01-JUN-2001; 2001WO-US017672.

2000US-0208538P. 2000US-0244989P. 2001US-00818875. 01-JUN-2000; 2 30-OCT-2000; 2 27-MAR-2001; 2

Wed Apr 21 12:58:21 2004

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(UYDE ) UNIV DELAWARE
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Kim J; Rice MC, Kmiec EB, Gamper HB, WPI; 2002-106307/14. New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 163; 220pp; English.

The invention relates to an oligomuclectide for targeted alteration of a genetic sequence, which comprises a single-stranded oligomuclectide features at DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligomuclectide. The chemical modifications of the oligomuclectide. The chemical modifications of plant genetic information of my two or more of those modifications. The oligomuclectides are useful for more of these modifications. The oligomuclectides are useful for directing repair or alteration of plant genetic information. The oligomuclectides are particularly useful for creating plants with desired of phenotypes, e.g. environmental or abiotic stress tolerance, improved nutritional value (e.g. altering amino acid content of plants or conferring amino acid content of plants or conferring mino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinone and sulphomylurea herbicide cresistance, indiadzolinone and sulphomylure are sestance, indiadzolinone and sulphomylure are production (e.g. increased starch or production, modified fatry acid content of e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatry acid content or the oligomuclectides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome content of plants of the invention

Sequence 17 BP; 1 A; 6 C; 7 G; 3 T; 0 U; 0 Other;

Gaps ; 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; rative 0; Mismatches 2; Indels Query Match Best Local Similarity 87.5' Matches 14; Conservative

Š a RESULT 466 ABK26640

ABK26640 standard; DNA; 17 BP. ABK26640;

09-APR-2002 (first entry)

Waxy starch production genome altering oligonucleotide #296.

Chromosomal genomic alteration; genome altering oligonucleotide; PCR; ss., o-methyl modification; LNA modification; phosphorothioate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abjocit stress tolerance; improved nutritional value, hygromycin, primary amino acid over production; herbicide resistance; glyphosate resistance; imidazolinone herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; triazine resistance; altered floral morphology; male sterile production; waxy starch; modified floral morphology; male sterile plant; albino mutant; modified fatty acid content; reduced plant; albino mutant; increased stearate production; reduced linolenic acid production; plant; increased stearate production; reduced linolenic acid production; photosynthetic process

Oryza glaberrima. Synthetic.

01-JUN-2001; 2001WO-US017672 WO200192512-A2 06-DEC-2001. 8\$6666666666666666666666665

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

(UYDE ) UNIV DELAWARE.

Kim J;

Gamper HB, Rice MC,

Kmiec EB,

WPI; 2002-106307/14.

New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 162; 220pp; English.

The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications of consist of o-methyl modification, an LNA modification, two or more consist of o-methyl modifications. The oligonucleotides are useful for more of these modifications. The oligonucleotides are useful for directing repair or alteration of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired oligonucleotides are particularly useful for creating plants with desired contentional value (e.g. altering amino acid content of plants or nutritional value (e.g. altering amino acid content of plants or conferring amino acid over production, herbicide resistance (e.g. cylphosate resistance, inidazolinone and sulphomylurea herbicide resistance, or diffied over production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content of e.g. increased starch or production of waxy starch), altered floral corphology (e.g. male-sterile plants) or modified fatty acid content of the oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome of altering oligonucleotide of the invention

Sequence 17 BP; 3 A; 7 C; 6 G; 1 T; 0 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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380 CCGCGACGACGCGCC 395 2 cadddadradddddd 17 ઠે 셤

ABV79109 standard; DNA; 17 BP RESULT 467 ABV79109 8xxxxxxxxxxxxxx

03-JAN-2003 (first entry) ABV79109;

Human HTPL scanning oligonucleotide SEQ ID 355.

Human, gene therapy, tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.

Homo sapiens

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30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
23-MAY-2001; 2001WO-US000669.
                      28-JAN-2002; 2002EP-00001167
                                                                                       WPI; 2002-676582/73.
                                                                      (AEOM-) AEOMICA INC.
    EP1229046-A2
             07-AUG-2002
                                                                               Zhan J;
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The present invention relates to human testis expressed Patched like correct invention relates to human testis expressed Patched like protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL has two isoforms, with a few single base pair differences between the two codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL codon in HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is considered to human chromosome loll21. HTPL and its coding sequence are useful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorders include disorders edexpession or activity of human HTPL. Such disorders include disorders of testis, or adrenal, adult and forcatal muscle or colon function. HTPL proteins and mucleic acids are clinically useful diagnostic markers and potental therapeutic agents for male infertility and cancer. The present oligonucleotide was used in an example from the invention Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL. Sequence 17 BP; 1 A; 8 C; 6 G; 2 T; 0 U; 0 Other; Example 2; Page 110; 718pp; English Similarity

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Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                       0; Gaps
3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                        Human HTPL scanning oligonucleotide SEQ ID 353.
                                                                         ABV79107 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                       03-JAN-2003 (first entry)
                                           14; Conservative
                                                                                                                                                                                                                                       ABV79107;
       Query Match
Best Local 9
                                                                                                                                                                RESULT 468
ABU79107
XX
AC ABU791
XX
XX
DT 03-JAN
XX
XX
DD Human;
XX
KW Human;
KW human;
KW male 9
KW proste
                                         Matches
                                                                               à
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Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic;

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The present invention relates to human testis expressed Patched like CC protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL protein (HTPL, see ABV78759 to ABV78762 and ABB98519 to ABB98520). HTPL concerns five single base pair changes introduces a premature stop codon in HTPL-8 (S for short) compared to HTPL-16. for long). HTPL codon in HTPL-8 (S for short) compared to HTPL-16. for long). HTPL codon in HTPL-8 (S for short) stone organisation with the Patched protein. The shares an overall structure organisation with the Patched protein. The chart of Patched, and is a potential tumour suppressor. HTPL is to that of Patched, and is a potential tumour suppressor. HTPL is important in regulating male germ cell development, and the HTPL gene was important in regulating male germ cell development, and the HTPL, and in cuspful for diagnosing a disorder caused by mutation in HTPL, and in therapy and manufacture of a medicament for treatment or prevention of therapy and manufacture of a medicament for treatment or prevention of the therapy and manufacture of a medicament for treatment or prevention of the HTPL. Such disorder sinclude disorders of testis, or adrenal, adult and therap and manrow, brain, kidney, lung, placenta, prostate, prostate, collingial muscle or colon function. HTPL proteins and mucleic acids are calinically useful diagnostic markers and potential therapeutic agents for male infermility and cancer. The present oligonucleotide was used in an encourted the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human ERG hammerhead ribozyme target sequence, Seq ID No 1084.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.28+02; rive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 1 A; 9 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 110; 718pp; English
                                                                                                                                                                                  30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
33-MAY-2001; 2001WS-02800669.
09-OCT-2001; 2001US-00864761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK18437 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 136 ccccccrcccccccicca 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 cccccrcccccccrccA 17
                                                                                                                                            28-JAN-2002; 2002EP-00001167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.0
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                   (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-676582/73
                        Homo sapiens
                                                               EP1229046-A2
                                                                                                        07-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK18437;
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ABK18437/C
D ABK1841
XX ABK1841
XX ABK1841
XX XX BB Human J
XX Human J
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09-APR-2002 (first entry)

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vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay Weber syndrome; leukaemia; se; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
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Jarvis T, Von Carlowitz I, Mcswiggen JA, Mclaughlin F, Randi AM;

Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

%\$GGGGGGGGGGGGGGGGGGGGGK%\$\$BBB\$\$B\$B\$B\$\$B\$\$B\$\$B\$\$

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's saccome, melanoma, conditions selected from cancer, lymphoma, Ewing's saccome, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, cumour angiogenesis, diabetic retinopathy, macular degeneration, surged with relating a patient having a condition associated with the level of ERG, by contacting a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more theraples under conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or conjunction with one or more of other theraples such as radiation or cell, by contacting (I) is useful for reducing ERG activity in a cell, by contacting (I) with RNM, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnostic of conditions and diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically carametic mucleic acids molecules which regulate expression of ERG, and crayeting genes that share homology with ERG gene or ERG fusion genes. ARM/1354-ABK2219 represent nucleic acids, including antiense and caramed Crayeting genes that share homology with regulate expression of ERG, and crayeting and call molecules which regulate expression of ERG, and crayeting genes are accompanied to the invention

Sequence 17 BP; 2 A; 4 C; 6 G; 0 T; 5 U; 0 Other;

Gaps . 0 Query Match

3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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RESULT 470 ABK18438/C ID ABK18438 standard, RNA, 17 BP. AC ABK18438,

Claim 4; Page 78; 149pp; English. 16-MAY-2001; 2001WO-US015866 16-MAY-2000; 2000US-00572021 (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LID. WPI; 2002-082995/11. WO200188124-A2 Homo sapiens. 22-NOV-2001 amberzyme.

Novel polymucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome. Human, hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; se; Osler-Weber-rendu syndrome; leukaemia; osteoporosis; DNAzyme; inozyme; Randi AM; Human ERG hammerhead ribozyme target sequence, Seq ID No 1085. Mcswiggen JA, Mclaughlin F, 16-MAY-2001; 2001WO-US015866. 16-MAY-2000; 2000US-00572021. Jarvis T, Von Carlowitz I, (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD. WPI; 2002-082995/11. WO200188124-A2. Homo sapiens. 22-NOV-2001. amberzyme.

The invention relates to a nucleic acid molecule (I) which down regulates corpression of an Ets-related gene (ERG). (I) is useful for treating capression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, conditions selected from the condition arthritis, psoriasis, verruca vulgaris, angiofibroma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tubercous sclerosis, port wine stains, Sturge Weber syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies to the treatment in the method comprises the use of one or more therapies conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or chardotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting (I) with RMM, in the presence of a divalent confusement as Mg2+. (I) is useful for diagnosis of conditions and cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as Mg2+. (I) is useful for diagnosis of conditions and cation such as hare homology with ERG gene or ERG fusion genes. Crargeting genes that share homology with ERG gene or ERG fusion genes. Crargeting genes that share homology with ERG gene or ERG fusion genes. Crargeting genes that share homology with ERG gene or ERG fusion genes. Crargeting genes that share homology with ERG gene or ERG fusion genes. Crargeting genes that share homology with ERG gene or ERG fusion genes. Crargeting genes that share homology with a call and mutations within diseased cells and mutations of expression of ERG, and crargeting genes and crargeting genes and conditions and conditions and conditions and conditions and cond

Claim 4; Page 78; 149pp; English.

Sequence 17 BP; 2 A; 4 C; 5 G; 0 T; 6 U; 0 Other;

; 0 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; cive 0; Mismatches 2; Indels . 0 Best Local Similarity 87.5 Matches 14; Conservative Query Match

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Сарв

408 393 GCCAAGAAGGTCTTCT

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16 gccaagaaggccarcr

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (G1, ABB83999), a sequence having 65% sequence identity to (S1), (S1), having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPsases as well as downstream components of the signal transduction pathway. (II) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) consetul or acused by altered expression of human POSHL1 including diagnosing and treating caused by altered expression of human POSHL1 including microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the constitution by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL -1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
                                                                                                                                                                                       Human; POSHL 1; SH3 domain; POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine; gene therapy; transgenic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 1747; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                      Human POSHL1 scanning oligonucleotide SEQ ID NO 1747.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 4 A; 7 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WS-US000669.
23-MAY-2001; 2001US-00864761.
                                     ABV91034 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                          28-JAN-2002; 2002EP-00001165
                                                                                                            23-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                            EP1239051-A2
                                                                                                                                                                                                                                                                                                                                                    11-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shannon M;
                                                                           ABV91034;
RESULT 47
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Matsumura Y, Moriya S, Nishida M;

Inoko H, Kagiya T, Ichihara T,

01-JUN-2000; 2000JP-00164798.

(NISN ) NISSHINBO IND INC. (SYST-) SYSTEM RES INC.

01-JUN-2001; 2001WO-JP004662.

WO200192572-A1. Homo sapiens

06-DEC-2001.

Human, human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.

Human HLA genotyping oligonucleotide SEQ ID NO 27

21-MAR-2002 (first entry)

ABL30538;

ABL30538 standard; DNA; 17 BP

ABL30538

339 CAGGGCCGCCTGCTCT 354

à

17 CAGGCCGGCTGTG

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                                                                         Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0
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Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 10; Page 98; 345pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    135 GCCCGCCTGCCGGTGG 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GACTGCCTGGCGGTGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACA09011 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-122074/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACA09011;
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ACA09011
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Gaps ö

sub-unit modulating amberzyme substrate #174

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; d-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; stomach cancer; colorectal cancer; pancreatic cancer; oesophageal cancer; head and neck cancer; varian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; gencitabine; radation therapy; inflammacory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; seepsis; transplant/graft rejection; reperfusion inlury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 07-DEC-1992; 18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

(STIN/) STINCHCOMB D T. (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G.

Draper KG; Stinchcomb DT, Mcswiggen J,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 54; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor crift shapes B NERBA), where (I) is an inozyme, decleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for creating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and cantisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cantisting resistant cancer. The method involves use of other drug charapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, colorectal, nectenoscial, fluorouracil carboplatin, edatrexate, colopsity, autoinmune disease, lupus, multiple sclend and antisense nucleic acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosus, asthma, Crohar disease, unducing molecules are also useful for treating inflammatory disease such as colorin, gene therapy applications, ischemial/repervices, ceptic, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic concept.

Sequence 17 BP; 5 A; 3 C; 8 G; 0 T; 1 U; 0 Other;

ö Gaps ö 3.0%; Score 12.8; DB 1; Length 17; ilarity 81.2%; Pred. No. 3.2e+02; Conservative 1; Mismatches 2; Indels Query Match Best Local Similarity Matches 13; Conserv

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286 CCAAGCTGGTGAAGGA 301
                 CCAGGCUGGGGAAGGA
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ACA06662 standard; RNA; 17 RESULT 474 ACA06662

ACA06662;

03-JUN-2003 (first entry)

NFKB sub-unit modulating inozyme substrate #481.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; escophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; head and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; pactitaxel; docetaxel; cisplatin; methotraxate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; crohn's disease; obesity; lacknemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

STIN/) STINCHCOMB D T. ٠, (MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G.

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 34; 72pp; English.

rhe invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NREM), where (I) is an inozyma, sinzyme, G-Claever or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotraxate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, gemcitabine or radiation therapy. The enzymatic and antisense nucleic

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Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
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acid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restemosis, asthma, Crohm's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; ocosophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; stomach cancer; ovarian cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumarcid arthritis; restenosis; Crohn; s disease; obstiv; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                          Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                       Sequence 17 BP; 0 A; 7 C; 9 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NFKB sub-unit modulating inozyme substrate #262.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-MAY-2001; 2001US-00864785.
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94US-00245466.
94US-00291932.
96US-00777916.
                                                                                                                                                                                                                                                                                                                                                                                                                             ACA06443 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                          305 GAGCCCCGGGGACCGC 320
                                                                                                                                                                                                                                                                                                                             16 dadcccccdddccccc 1
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MCSW/) MCSWIGGEN J.
DRAP/) DRAPER K G.
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15-AUG-1994;
23-DEC-1996;
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ACA06443/c
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configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. treating a patient having a condition associated with the level of REL-A. The presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, displant, methotrexate, colorectherapy including paclitaxel, docetaxel, displant, edatrexate, gencitables are also useful for treating inflammatory disease such as the manual disease, lung, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury rejection, alexamen (CNS) and myocardial), glomerulonephritis, espenies alrawy inflammatory bowel disease or infection, allergic alrawy inflammation, inflammatory bowel disease or infection, and inflammation, inflammatic, septements the substrate of a novel enzymatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cospapageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; methorreate; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; cyclophosphamide; docetaxel; docetaxel; cisplatin; methorrexate; cyclophosphamide; docorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoind arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
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3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 2 A; 10 C; 3 G; 0 T; 2 U; 0 Other;
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94US-00245466.
94US-00291932.
96US-00777916.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   142 TGGCGGTGGAGGCCGG 157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid molecule
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15-AUG-1994;
23-DEC-1996;
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Stinchcomb DT, Mcswiggen J, Draper KG,

(DRAP/) DRAPER K G.

WPI; 2003-340953/32.

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regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg-2+, the enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophagel, stoomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, completable or radiation televal, action antisense nucleic genitable or radiation therapy. The enzymatic and antisense nucleic caid molecules are also useful for treating inflammatory disease such as molecules are also useful for treating inflammatory disease, cobesity, autonimume disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischeemia/reperfusion injury content nervous system (CNS) and myocardial), glomerulonephritis, cepties, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic connection. This sequence represents the substrate of a novel enzymatic
Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                         describes an enzymatic nucleic acid molecule (I) which down
                                                                                                                                                                                                                                                                                           Claim 3; Page 34; 72pp; English.
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Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels Sequence 17 BP; 0 A; 6 C; 9 G; 0 T; 2 U; 0 Other; 305 GAGCCCCGGGGACCGC 320

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0; Gaps

ACA06586 standard; RNA; 17 BP. 17 chácccccccccc 2 RESULT 477 4CA06586/c ð 셤

03-JUN-2003 (first entry) ACA06586; 

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; dung cancer; prostate cancer; colosectal cancer; brain cancer; colosectal cancer; brain cancer; bronded cancer; become cancer; cancer; cervical cancer; stomach cancer; bladder cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paolitaxal; docetaxel; cisplatin; methorraxate; cyclophosphanide; docetaxel; cisplatin; methorraxate; cyclophosphanide; docetaxel; cisplatin; detrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; pheumatoid arthritis; reserobals; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss. NFKB sub-unit modulating inozyme substrate #405.

Draper KG; 92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. Mcswiggen J, 23-MAY-2001; 2001US-00864785. STINCHCOMB D T. MCSWIGGEN J. DRAPER K G. JS2002177568-A1 Stinchcomb DT, 5-AUG-1994; 23-DEC-1996; 07-DEC-1992; 18-MAY-1994; 28-NOV-2002 (STIN/) (DRAP/)

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases. WPI; 2003-340953/32.

Claim 3; Page 33; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, 2inzwe, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat configuration associated with the level of KEL-A creating a patient having a condition associated with the level of KEL-A. (I) is useful for cleaving KNA comprising a sequence of KEL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and anticense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, carvical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chenotherapy including paclitaxel, docetaxel, cisplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, colection, and arthritis, restenosis, asthma, Crohn's disease such as entenosis, restenosis, sethemia/reperfusion injuny, colection, and arthritis, restenosis, suchma, crohn's disease, diabetes, obesity, autoimmume disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injuny, seppis, allergic airway inflammatory bowel disease or sepsis, allergic airway inflammation, inflammatory bowel disease or infection, and an expense or sepsis, allergic airway inflammation, inflammatory bowel disease or infection. The sequence represents the substrate of a novel enzymatic

Sequence 17 BP; 1 A; 7 C; 6 G; 0 T; 3 U; 0 Other;

Gapa ö Query Match
3.0%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 3.2e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

266 GCACCTGGAGCAGGGC 281

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ACA09010 standard; RNA; 17 BP 17 dcagcracadcaddd 2 ACA09010; RESULT 478 ACA09010 d 

NFKB sub-unit modulating amberzyme substrate #173.

(first entry)

03-JUN-2003

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

Homo sapiens

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G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; broadged cancer; brain cancer; carcer; carcer; carcer; cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restensis; Crohn's disease; obseity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
                                                                                                                                              Homo sapiens.
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US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132, 94US-00245466, 94US-00291932, 96US-00777916, 07-DEC-1992;

18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

STIN/) STINCHCOMB D T. MCSWIGGEN J.

DRAPER K G. Stinchcomb DT, DRAP/)

Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 54; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (1) which down regulates expression of a sequence encoding a subunit of nuclear factor adapa B (NFXB), where (1) is an inozyme, zinzyme, G-cleaver or amberzyme cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and nack, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paciltaxel, docetaxel, cisplantin, methotrexate, gemiciabine or radiation therapy. The enzymatic and antisense nucleic scid molecules are also useful for treating inflammatory disease such as theumatoid arthritis, resenosis, asthma, crohn's disease, diabetes, cheumatoid disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, isohaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sequence represents the substrate of a novel enzymatic nucleic acid molecule

Sequence 17 BP; 4 A; 4 C; 8 G; 0 T; 1 U; 0 Other;

Gaps . Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 81.2%; Pred. No. 3.2e+02; Matches 13; Conservative 1; Mismatches 2; Indels

2 CCAGGCUGGGGAAGGA 17

RESULT 479 ADA99410

ADA99410 standard; DNA; 17

ADA99410;

20-NOV-2003 (first entry)

Human MDZ3 scanning oligonucleotide SEQ ID 399.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MD23; MD24; MD212; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181 (AEOM-) AEOMICA INC. Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 399; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ3 is cancoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 1671.2 and MDZ12 is encoded at chromosome 1671.2 and MDZ12 is encoded at chromosome 1671.2 and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The mucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids and proteins are also useful for disquesting cancerize gross caids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as 

Sequence 17 BP; 3 A; 7 C; 1 G; 6 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 3.2e+02; Artive 0; Mismatches 2; Indels 14; Conservative Query Match Best Local Similarity Matches

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RESULT 480 ABZ61658

ABZ61658 standard; RNA; 17 BP.

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286 CCAAGCTGGTGAAGGA 301

ABZ61658;

Page 233

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14; Conservative
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Roberts B;
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DRAPER K.
ROBERTS E.
                                                                                                              Hepatitis C virus.
                                                                                                                                                                                           WO200281494-A1
                                                                                                                                                                                                                                                                     17-0CT-2002
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Draper K,
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AC ACD587
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                   Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                              Human H-Ras DNAzyme target #449.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 481
ACD58640/c
ID ACD58640 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                287 CAAGCTGGTGAAGGAC 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-2002; 2002WO-US016840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200297114-A2.
                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-DEC-2002
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HHV) RNB. The nucleic acid molecules include antisense and enzymatic nucleic acids such as harmerhead ribozymes, DNBxzymes, CC are nucleic acid decoup molecules and aptemers that DNBxzymes, Also disclosed are nucleic acid decoy molecules and aptemers that bind to HBV reverse transcriptase primer sequences, as well transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNB. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds compounds in modulate the expression and/or replication of HCV. The compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carpinession ench as cirrhosis, liver failure, and hepatocellular of the HCV DNAzme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ij
liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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87.5%; Pred. No. 3.2e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Morrissey D, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 1 A; 8 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 250; 387pp; English.
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08-JUN-2001, 2001US-00877478.
08-JUN-2001, 2001US-0296876P.
24-OCT-2001, 2001US-0335059P.
05-DEC-2001, 2001US-0337055P.
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BLATT L.
MACEGJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACD58724 standard; RNA; 17
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24-SEP-2003 (first entry)

Wed Apr 21 12:58:21 2004

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Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus, HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; inzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative, disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J, Morrissey D, Pavco P,
HCV DNAzyme substrate sequence #974.
                                                                                                                                                                                                                                                                                                                                                                             26-MAR-2002; 2002WO-US009187.
                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2001; 2001US-00817879.
                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIBOZYME PHARM INC.
BLAIT 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ROBERTS E.
                                                                                                                                                                                                                                          Hepatitis C virus
                                                                                                                                                                                                                                                                                      WO200281494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Blatt L, Mo
                                                                                                                                                                                                                                                                                                                                17-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
(DRAP/)
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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus Claim 1; Page 251; 387pp; English. nfection.

WPI; 2003-229207/22.

The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as harmarchead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed are nucleic acid decoy molecules and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligomucleotides that specifically bind the Enhancer I region of HBV Genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV inverse.

Sequence 17 BP; 2 A; 7 C; 5 G; 0 T; 3 U; 0 Other;

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Gaps
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0
3.0%; Score 12.8; DB 1; Length 17; ilarity 87.5%; Pred. No. 3.2e+02; Conservative 0; Mismatches 2; Indels
  Query Match
Best Local Similarity
Matches 14; Conserv
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New human sodium-hydrogen exchanger like protein 1 (NHELP1), useful as passive replacement therapy or as a vaccine for treating or preventing disorders associated with aberrant expression or activity of human
                                                                                                                                       ss; gene therapy; vaccine; sodium/hydrogen exchanger like protein; NHELP1; passive replacement therapy; vaccine; diagnosis.
                                                                                                                    Human Na/H exchanger-like protein 1 gene oligonucleotide #702
                                                                                                                                                                                                                                           30-JAN-2001; 2001WO-US000666.
23-MAY-2001; 2001US-00864761.
21-DEC-2001; 2001US-0343331P.
310 CCGGGGACCGCGTGCT 325
                                                                                                                                                                                                                           25-JAN-2002; 2002EP-00001160
                 16 cccccccccci
                                                              ADC04255 standard; DNA; 17
                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                     WPI; 2003-302724/30.
                                                                                                                                                                                                                                                                                 (AEOM-) AEOMICA INC
                                                                                                                                                                     Homo sapiens,
                                                                                                                                                                                      EP1273660-A2.
                                                                                                    18-DEC-2003
                                                                                                                                                                                                        08-JAN-2003.
                                                                                  ADC04255;
                                                                                                                                                                                                                                                                                                                                                                   NHELP1.
                                                                                                                                                                                                                                                                                                  Gu Y;
                                              RESULT
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The invention relates to a nucleic acid molecule which encodes a Na+/H+ exchanger like protein (NHELPI). The NHELPI nucleic acid molecule, NHELPI polypeptide, an antibody against the protein or its antighen-binding fragment is useful in therapy. The NHELPI nucleic acid molecule, NHELPI polypeptide and an agonist are particularly useful for manufacturing a medicament for treating or preventing a disorder associated with antigen-binding fragment, and an antagonist, are useful for manufacturing a medicament for treating or preventing a disorder associated with increased expression or activity of human NHELPI. The NHELPI nucleic acid or protein is useful as passive replacement therapy, as a vaccine, or in diagnostic methods. This sequence corresponds to a 17-mer oligonucleotide spanning the sequence of the human NHELPI gene (ADC03514).

Example 2; SEQ ID NO 742; 468pp; English.

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Gaps
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                                             Query Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
Sequence 17 BP; 2 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                        39 GAAGATGGCCACCACT 54
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ADC04256/c ID ADC04256 standard; DNA; 17 BP. XX RESULT 484

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GAAAATGGCCAGCACT

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The invention relates to a nucleic acid molecule which encodes a Na+/H+ exchanger like protein (NHELP1). The NHELP1 nucleic acid molecule, NHELP1 polypeptide, an antibody against the protein of standing fragment is useful in therapy. The NHELP1 nucleic acid molecule, NHELP1 polypeptide and an agonist are particularly useful for manufacturing medicament for treating or preventing a disorder associated with decreased expression or activity of human NHELP1. The antibody or its antigen-binding fragment, and an antagonist, are useful for manufacturing a medicament for treating or preventing a disorder associated with increased expression or activity of human NHELP1. The NHELP1 nucleic acid or protein is useful as passive replacement therapy, as a vaccine, or in diagnostic methods. This sequence corresponds to a 17-mer oligonucleotide spanning the sequence of the human NHELP1 gene (ADC03514).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plant growth; plant growth trait modulation; Brassicaceae; Arabidopsis; Brassica; Zea; Oryza; Triticum; Hordeum; Lolium; Sorghum; Glycine; Medicago; Helianthus; Lactuca; Beta; Vitis; Solanum; Lycopersicon; Capsicum; Gossypium; Hevea; Linum; Prunus; Citrus; Populus; Pinus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New human sodium-hydrogen exchanger like protein 1 (NHELP1), useful as passive replacement therapy or as a vaccine for treating or preventing disorders associated with aberrant expression or activity of human
                                                                                                            ss; gene therapy; vaccine; sodium/hydrogen exchanger like protein; NHELP1; passive replacement therapy; vaccine; diagnosis.
                                                                         Human Na/H exchanger-like protein 1 gene oligonucleotide #703.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 3.0%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 3.2e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Plant growth associated polynucleotide seg id 203.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 2 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 743; 468pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE25228 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                               30-JAN-2001; 2001WO-US000666.
23-MAY-2001; 2001US-00864761.
21-DEC-2001; 2001US-0343331P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39 GAAGATGGCCACCACT 54
                                                                                                                                                                                                                                                                                           25-JAN-2002; 2002EP-00001160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 GAAAATGGCCAGCACT 1
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                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-302724/30.
                                                                                                                                                                                                                                                                                                                                                                                                                (AEOM-) AEOMICA INC.
                                     18-DEC-2003
                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                EP1273660-A2
                                                                                                                                                                                                                                                      08-JAN-2003.
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  ADC04256;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gu Y;
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ADE25228
MAKAKEKAKAKA
MAKAKEKAKAKA
MAKAKEKAKAKA
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The invention describes an isolated or recombinant polypeptide (I) comprising a sequence: (a) comprising 1 of 30 sequences (S1), as given in the specification, or a conservative variant; (b) encoded by 1 of 30 sequences (S2), as given in the specification, or a conservative variant; (c) encoded by a sequence that hybridises under stringent conditions to activity of (I) is modulated to modulate a plant growth trait in a city Arabidopsis, Brassica, Zea, pricticum, Hordeum, Lollum, Sorghum, Glycine, Medicago, Helianthus, Lactuca, Beta, Vitis, Solanum, Lycopersicon, Capsicum, Gossypium, Hevea, Linum, Prunus, Citrus, Populus, Brus, or Quercus, A membrod is used to detect genes for a plant growth trait. This sequence represents a polynucleotide isolated from the plant growth associated genes of the invention that can be used a a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Probe, component B; promoter; human; signal peptide; primer; RACE; low molecular weight protein; urine; TGF-alpha; receptor; amplify; inflammation; coagulation; tumour; anglogenesis; ss.
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87.5%; Pred. No. 3.2e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                  Example 2; SEQ ID NO 203; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer, probe or genetic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Component B gene primer, CKCB2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ87873 standard; DNA; 18 BP.
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Local Similarity 87.5
Les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
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27-JUL-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ87873;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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AAQ87873/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       $X8X#X#X#X#X#X#X#X#X#X#X#X#X
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Gaps

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New isolated or recombinant polypeptide for use in modulating a plant growth trait in a flowering plant e.g. in Arabidopsis, Brassica, Zea, or

Bowen BA, Haudenschild CD, Buckler ES

WPI; 2003-803305/75.

Oryza.

(LYNX-) LYNX THERAPEUTICS INC.

09-JAN-2002; 2002US-0347288P. 07-JAN-2003; 2003US-00338777

US2003188343-A1 Magnoliophyta.

02-OCT-2003.

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22-DEC-1992; Query Match Best Local S Sirna A; RESULT 487 AAA58496 쉱 ò

285 ACCAAGCTGGTGAAGG 300 14; Conservative

17 Accacecreereaces 2

AAA58496 standard; DNA; 18 BP 20-OCT-2000 (first entry) AAA58496; 

PCR primer used to amplify bleomycin (BLM) gene cluster ORF19.

BIM gene cluster; bleomycin gene cluster; polyketide metabolite; bleomycin; bleomycin analogue; holo-carrier protein; thiazolidine; thiazoline; bithiazoline; microbial metabolite; sugar; PCR primer;

WO200040704-A1.

06-JAN-2000; 2000WO-US000445

06-JAN-1999; 05-FEB-1999; 05-JAN-2000; 2

Shen B,

New bleomycin gene cluster components useful for peptide and/or polyketide metabolites, especially bleomycin, production and for chemically modifying biological molecules.

Disclosure; Page 22; 162pp; English

PCR primers AAAS8474-A58541 were used to amplify open reading frames (ORFs) 8 to 41 of the BLM (Bleomycin) gene cluster. The proteins encoded by the gene cluster are useful for producing peptides and/or polyketide metabolites, especially bleomycin or bleomycin analogues. They are also useful for chemically modifying biological molecules to produce branched reacted with an apo-carrier protein and coenzyme A to produce a holocarrier protein. The BLM gene cluster or catalytic domains can be used individually or collectively to produce thiazolidine, thiazoline, bithiazoline and bithiazoline-containing microbial metabolites. The BLM gene cluster may also be used to produce sugars

Sequence 18 BP; 3 A; 5 C; 8 G; 2 T; 0 U; 0 Other;

0; Gaps Query Match
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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294 GTGAAGGACCTGAGCC 309 1 GTGAAGGACCTCGGCC 16 셤 ઠ

AAH40454 standard; DNA; 18 BP (first entry) 14-AUG-2001 AAH40454; RESULT 488 AAH40454/c

SNP specific lower PCR primer SEQ ID 3250.

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPB; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; zheumaroid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.

Homo sapiens.

WO200129262-A2

26-APR-2001.

99US-0160096P. 13-OCT-2000; 2000WO-US028436. 15-OCT-1999;

(ORCH-) ORCHID BIOSCIENCES INC. Piccult-Newburg L, Pohl M;

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic acid sample. WPI; 2001-290930/30. 

Claim 1, Page 66; 83pp; English

(REGC ) UNIV CALIFORNIA Du L'

Sanchez C,

Edwards DJ; Chen M, WPI; 2000-465974/40.

&XTXEXXFFFFXXXXOOOOOOOOO

92IT-RM000919

(ISTF ) ARS APPLIED RES SYST HOLDING NV.

WPI; 1994-234696/28.

New protein, component B, isolated from urine - with antiinflammatory, anticoagulant and anti-tumour activities, also related nucleic acid, vectors and transformed cells.

Example 4; Page 28; 55pp; English.

The sequences given in AAQ87870-75 are primers which were used in the amplification of the component B cDNA. These primers were used in the rapid amplification of cDNA ends (RACE) and are targetted to various credions of the gene including exon 2 and the poly-A tail. The component B gene contains three exons and two introms. Exon 1 is 84 bp and contains cas bases of untranslated mRNA. It encodes 19 amino acids of the putative signal peptide and is separated from exon 2 by an intron of 410 bp. Exon 2 is 120 bp and codes for 3 amino acids of the putative signal sequence and 37 amino acids of the mature protein. It is separated from exon 3 by an intron of abbout 550 bp. Exon 3 is 326 bp and encodes the C-terminal contains a poly-A signal 14 bp upstream of the 3 processing site.

Component B is a low molecular weight protein which may be isolated from thum unine by adsorption at acid pH on kaclin, then extraction with column hydroxide. It inhibits binding of TGP-alpha to its receptor, and so has antiinflammatory, anticoagulant and/or antitumour activities. It may also be used to treat conditions associated with altered levels of TGF-alpha, eg. behavioural or hormonal disturbances and angiogenesis. See 25 mAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)

Sequence 18 BP; 3 A; 7 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ; 0 3.0%; Score 12.8; DB 1; Length 18; Similarity 87.5%; Pred. No. 3.6e+02; Conservative 0; Mismatches 2; Indels

88

Streptomyces verticillus,

13-JUL-2000.

99US-0115435P. 99US-0118848P. 2000US-00477962.

Page 237

Wed Apr 21 12:58:21 2004

Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide polymorphisms SNPs. The present invention of sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the includes kits for determining the presence or absence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a SC primers are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The performing a single-nucleotide primer extension reaction. The cligonucleotides are useful for determining the presence, absence or dentity of a SNP and for genotyping nucleic acid sample by performing a single-nucleotide primer extension reaction. The cligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic traits include diseases e.g. caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinemia, diabetes insipidus, Lesch Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial clisaase of which a component is or may be genetic such as autoimmune disease including, rheumatoid arthritis, multiple sclerosis, including, rheumatoid arthritis, multiple sclerosis, incroorganism. The method is also useful in forensic investigations and containing DNA sequence represents a PCR primer specific cor a numan SNP containing DNA sequence

Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 U; 0 Other;

0; Gaps / Match 3.0%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 3.6e+02; nes 14; Conservative 0; Mismatches 2; Indels Query Match Best Loca Matches ે

299 GGACCTGAGCCCCGGG 314 18 dérecrahadeceades

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ABL40174 standard; DNA; 18 BP ABL40174; RESULT 489 ABL40174 

(first entry) 21-MAY-2002 Mouse reelin protein CR-50 epitope region PCR primer SEQ ID NO:11.

Mouse; realin protein CR-50 epitope region; elucidation; neuron; cerebral disturbance; reelin protein; neuroprotective; PCR primer; ss.

Mus musculus.

JP2002017361-A.

22-JAN-2002.

04-JUL-2000; 2000JP-00202801.

04-JUL-2000; 2000JP-00202801

(RIKE ) RIKEN KK.

WPI; 2002-221707/28.

Reelin protein CR-50 epitope region, useful for diagnosis and treatment of cerebral disturbance.

Example 2; Page 7; 16pp; Japanese.

The present invention describes the mouse realin protein CR-50 epitope region, which contains the CR-50 antibody recognition site and is free from F-spondin domains and repetitive sites. Also described are: (1) an expression vector comprising a polymucleotide encoding a realin protein epitope region; (2) host cells with transfected the expression vector;

(3) polypeptides prepared by culture of the host cells; and (4) polymucleotides comprising the 351 base sequence given in ABL40165 which encodes the 117 amino acid sequence given in ABB06244; and (5) use of the polymucleotide for diagnosis and/or treatment of diseases caused by abnormal positioning of neural cells, and stimulation of association of reelin protein. The mouse reelin protein CR-50 epicope region has neuroprotective activity, and can be used in the diagnosis and treatment of cerebral disturbance due to an abnormal reelin gene and positioning of neurons. The present sequence represents a PCR primer for the mouse reelin protein CR-50 epitope region, which is used in an example from the present invention 88888888888888

Sequence 18 BP; 4 A; 5 C; 6 G; 3 T; 0 U; 0 Other;

Gaps 3.0%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 3.6e+02; Aztive 0; Mismatches 2; Indels 0; Local Similarity 87.5 es 14; Conservative Query Match Matches

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RESULT 490 ABK27438/c

ABK27438 standard; DNA; 18

ABK27438;

09-APR-2002 (first entry)

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Colon cancer associated cDNA CATX-7, 5' PCR primer.

Human; colon cancer; tumour; abnormal cell growth; melanoma; cervical cancer; colorectal adenocarcinoma; Wilms' tumour; leukaemia; lymphoma; antisense therapy; CATX; probe; primer; ss.

Homo sapiens.

WO200111047-A2.

15-FEB-2001.

08-AUG-2000; 2000WO-US021606.

09-AUG-1999; 99US-0147933P.

(FARB ) BAYER CORP.

Bowman BM, Wang K;

WPI; 2002-121548/16.

New isolated nucleic acid involved in growth regulation in human colonic epithelial cells, termed CATX, for diagnosing and treating abnormal cell growth, and for use as a probe/primer for detecting tumors.

Example; Page 88; 130pp; English.

The invention relates to an isolated nucleic acid (I) involved in growth regulation in human colonic epithelial cells, termed CATX. (I) is useful as a probe-primer for detecting tumours, preferably colon cancer. The nucleic acid, encoded polypeptide and antibody are useful in diagnosis and treatment of abnormal cell growth (such as cervical cancer, mealsonmas, colorectal adenocarcinomas, Wilms' tumour, leukaemias and lymphomas), in screening assays for the treatment of abnormal cell growth, for raising antibodies, and to screen for peptide analogues and growth, for raising antibodies, and to screen for peptide analogues and printaghists. (I) is useful as a biomarker for human tumour cells, e.g., colon cancer cells, for generating probes and primers designed for identifying and/or cloning homologues in other cell types, in antisense brange of development, so that premalignant cells can be identified prior to their spreading throughout the human body. (I) allows early detection 

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Page 238
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of potentially cancerous conditions, and treatment of the cancerous conditions prior to spread of the cancer cells throughout the body, or prior to development of an irreversible cancerous condition. ABXC7426-ABX27469 represent human colon cancer associated coding sequences and primers of the invention

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Sequence 18 BP; 3 A; 5 C; 9 G; 1 T; 0 U; 0 Other;

Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels 338 CCAGGGCCGGCTGCTC 353 à

18 CCAGGGCTGGCTCTC 3 셤

ABK27436 standard; DNA; 18 BP ABK27436; 

Colon cancer associated cDNA CATX-6, 5' PCR primer. 09-APR-2002 (first entry)

Human; colon cancer; tumour; abnormal cell growth; melanoma; cervical cancer; colorectal adenocarcinoma; Wilms' tumour; leukaemia; lymphoma; antisense therapy; CATX; probe; primer; ss.

Homo sapiens.

WO200111047-A2.

15-FEB-2001.

08-AUG-2000; 2000WO-US021606

99US-0147933P. 09-AUG-1999;

(FARB ) BAYER CORP.

Bowman BM, Wang K;

WPI; 2002-121548/16.

New isolated nucleic acid involved in growth regulation in human colonic epithelial cells, termed CATX, for diagnosing and treating abnormal cell growth, and for use as a probe/primer for detecting tumors.

Example; Page 88; 130pp; English.

The invention relates to an isolated nucleic acid (I) involved in growth regulation in human colonic epithelial cells, termed CATX. (I) is useful as a probe/primer for detecting tumours, preferably colon cancer. The nucleic acid encoded polypeptide and antibody are useful in diagnosis and treatment of abnormal cell growth (such as cervical cancer.) The cancer plants of an inscreening assays for the treatment of abnormal cell growth, for raising antibodies, and to screen for peptide analogues and antagonists. (I) is useful as a biomarker for human tumour cells, e.g., colon cancer cells, for generating probes and primers designed for identifying and/or cloning homologues in other cell types, in antisense therapy, and in tissue profiling. (I) identifies cancer cells at an early category and in tissue profiling. (I) identifies cancer cells at an early category are spreading throughout the human body. (I) allows early detection to conditions prior to spread of the cancer cells throughout the body, or prior to development of an irreversible cancerus condition. ABKZ7465 primers of the invention

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                                   Query Match
3.0%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
Sequence 18 BP; 3 A; 5 C; 9 G; 1 T; 0 U; 0 Other;
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'432/c ABK27432 standard; DNA; 18 RESULT 492 ABK27432/c

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09-APR-2002 (first entry) ABK27432;

Colon cancer associated cDNA CATX-4, 5' PCR primer.

Human, colon cancer; tumour, abnormal cell growth; melanoma; cervical cancer; colorectal adenocarcinoma; Wilms' tumour; leukaemia; lymphoma; antisense therapy; CATX; probe; primer; 88.

Homo sapiens

WO200111047-A2.

15-FEB-2001

08-AUG-2000; 2000WO-US021606.

99US-0147933P 09-AUG-1999;

(FARB ) BAYER CORP.

Bowman BM, Wang K;

WPI; 2002-121548/16.

New isolated nucleic acid involved in growth regulation in human colonic epithelial cells, termed CATX, for diagnosing and treating abnormal cell growth, and for use as a probe/primer for detecting tumors.

Example, Page 87; 130pp; English.

The invention relates to an isolated mucleic acid (I) involved in growth regulation in human colonic epithelial cells, termed CATX. (I) is useful as a probe/primer for detecting tumours, preferably colon cancer. The nucleic acid, encoded polypeptide and antibody are useful in diagnosis and treatment of abnormal cell growth stumour, leukaemias and antanomas, colorectal adenocarcinomas, wilms, tumour, leukaemias and lymphonas), in screening assays for the treatment of abnormal cell growth, for raising antibodies, and to screen for peptide analogues and growth, for raising antibodies, and to screen for peptide analogues and antagonists. (I) is useful as a biomarker for human tumour cells, e.g., colon cancer cells, for generating probes and primers designed for identifying and/or cloning homologues in other cell types, in antisense therapy, and in tissue profiling. (I) identifies cancer cells at an early crappy and in tissue profiling. (I) identifies cancer cells at an early crappy and in colon cancer cells and concercing conditions prior to spread of the cancer cells can be identified prior to their spreading throughout the human body. (I) allows early detection of potentially cancerous conditions and treatment of the cancerous conditions prior to gread of the cancer cells throughout the body. Or prior to development of an irreversible cancerous condition. ABK27426-CC primers of the invention 

Sequence 18 BP; 3 A; 5 C; 9 G; 1 T; 0 U; 0 Other;

Gaps ö Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels

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Monoclonal antibody; fusion protein; antigen; cell surface; receptor; ss.
                                                                                                                                                                                                            A protein fused with a monoclonal antibody against an antigen present on
                                                                                     Monoclonal antibody related oligonucleotide.
                                                                                                                                                                                                                                Example; Page 6; 24pp; Japanese.
                                             ABA94181 standard; DNA; 18 BP.
338 CCAGGGCCGGCTGCTC 353
                                                                                                                                                                    29-MAY-2000; 2000JP-00158575.
                                                                                                                                                       29-MAY-2000; 2000JP-00158575.
            18 ccadederectrers
                                                                        (first entry)
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                                                                                                                             JP2001333780-A.
                                                                                                                                                                                                                   cell surfaces
                                                                        09-MAY-2002
                                                                                                                                          04-DEC-2001
                                                                                                                Synthetic.
                                                          ABA94181;
                                       ABA94181
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The present invention describes a protein which is fused with a monoclonal antibody against an antigen present on cell surface and which can transfer a gene by combining with the gene and containing a human type single-stranded monoclonal antibody and a peptide which is the combining site for the gene. Also described is a complex of a monoclonal antibody-fused protein which is a complex of monoclonal antibody-fused protein and a method for the preparation of a monoclonal antibody-fused protein against a receptor present on cell surface in antibody-fused protein against a receptor present on cell surface in said monoclonal antibody against a receptor present on cell surface is used as the template to amplify a single-stranded antibody gene of a course type monoclonal antibody by PCR; (2) the framework portion of the mouse type monoclonal antibody is converted to prepare a single-stranded antibody gene of a human type monoclonal antibody to prepare a human type gene of the human type monoclonal antibody to prepare a human type single-stranded immunoporter gene; and (4) the human type single-stranded immunoporter gene; and (4) the human type single-stranded immunoporter gene; and (4) the human type single-stranded immunoporter gene; and is a microbacting the above complex of monoclonal antibody-fused protein of the human type single-stranded immunoporter. Also described is a method for introducing the above complex of monoclonal antibody-fused protein of the through a cell surface receptor. The method is used for the preparation of surface. The present sequence represents an oligonucleotide which is used in an example from the present invention
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Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
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Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.6e+02; Matches 14; Conservative 0; Mismatches 2; Indels GGTGCACCTGGAGCAG 278 263

GGTGCAGCTGCAGCAG 18 m

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The present invention relates to human chondroitin 6-sulfotransferase (C6ST) proteins and polynucleotides encoding such proteins. Sequences of the invention are useful in the molecular study of human extracellular matrix, for studying the biological functions of chondroitin 6-sulphate (C6S), in screening test for detecting C6ST polymorphs, for ascertaining and evaluating the role C6ST plays in atheroscierosis and for identifying potential therapeutics, i.e., inhibitors of enzyme or modulators of gene expression. The present DNA sequence is a PCR primer which is used for amplifying human C6ST gene
                                                                                                                                        Human, chondroitin 6-sulfotransferase, C6ST, chondroitin 6-sulphate, C6S, biological function; extracellular matrix; atherosclerosis; therapeutic; gene expression; enzyme; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel recombinant human chondroitin 6-sulfotransferase polynucleotide segment, useful in molecular study of human extracellular matrix, and for studying biological functions of chondroitin 6-sulfate.
                                                                                                           Human C6ST gene amplifying 5' PCR primer #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Col 17; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                        (UYJE-) UNIV JEFFERSON THOMAS
               AAD41288 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                         98US-00015188.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-535977/57.
                                                                                                                                                                                                                                                                                                                                                                                                                       Williams KJ,
                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                         29-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                        31-MAR-1997;
02-JUL-1997;
                                                                          30-OCT-2002
                                                                                                                                                                                                                                                                         04-JUN-2002.
                                            AAD41288;
AAD41288
ID AAD4
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Gaps ö Query Match 3.0%; Score 12.8; DB 1; Length 18; Best Local Similarity 87.5%; Pred. No. 3.66+02; Matches 14; Conservative 0; Mismatches 2; Indels

Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

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293 GGTGAAGGACCTGAGC 308 cercaaccaccided 17

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AAD24955 standard; DNA; 18 BP.

AAD24955;

Human beta IG-H3 promoter DNA amplifying antisense PCR primer. 12-MAR-2002 (first entry) 

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Gaps . 0

Human; growth inhibitory gene; retinoid; retinoic acid response element; RARE site; therapy; promyelocytic leukaemia; cancer chemoprevention; cytostatic; secreted cell adhesion protein beta IG-H3 promoter; PCR primer; ss.

Homo sapiens

RESULT 494

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Tang

Cheng Y,

Shen B,

WO200192578-A2

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Leinamycin biosynthesis gene cluster; Lmn; open reading frame; ORF; anti-tumour antibiotic; broad spectrum antimicrobial activity; darm-positive; Gram-negative bacteria; chemical modification; metabolite; apo-carrier protein; holo-carrier protein; tumour; polyketide; hybrid polypeptide/polyketide metabolite; Inm production; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer #1 for S. atroolivaceus leinamycin gene cluster ORF lnmQ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABX34392 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       104 TGACCGCGACCGCAGC 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptomyces atroolivaceus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 87.5'
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   d
                                                                                                                                                                                                                                                                           The patent discloses growth inhibitory genes induced by retinoids. The invention also relates to recombinant expression constructs that express a reporter gene under the transcriptional control of a promoter for a gene which is expressed by retinoid induction. The promoter does not contain a retinoic acid response elements (RARE) site. The invention that her tealates to reagents and methods for identifying compounds other than retinoids that modulate the expression of cellular genes. These compounds are useful for treating cancers such as promyelocytic leukaemia and cancer chemoprevention. The present DNA sequence is a PCR primer which is used for amplifying human secreted cell adhesion protein beta IG-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leinamycin biosynthesis gene cluster; Lmn; open reading frame; ORF; anti-tumour antibiotic; broad spectrum antimicrobial activity; dram-negative bacteria; chemical modification; metabolite; apo-carrier protein; holo-carrier protein; tumour; polyketide; hybrid polypeptide/polyketide metabolite; Inm production; cytostatic;
                                                                                                                                                                                                  Expression construct encoding cellular genes, under control of a promoter regulated by retinoids and cells comprising the construct for identifying compounds that induce expression of the genes useful in treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9.0%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 3.6e+02; los 14; Conservative 0; Mismatches 2; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer #1 for S. atroolivaceus leinamycin gene cluster ORF lnmM.
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                                                                                                                                                                                                                                                        Example 3; Page 27; 64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              366 CTCACTTTCCTGGACC 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX34384 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (REGC ) UNIV CALIFORNIA.
(KYOW ) KYOWA HAKKO KOGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAR-2002; 2002WO-US008937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-MAR-2001; 2001US-0278935P
                                                                 25-MAY-2001; 2001WO-US017161.
                                                                                            26-MAY-2000; 2000US-0207535P.
                                                                                                                                                 Dokmanovic M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      crcacrrccrreage 3
                                                                                                                     (UNII ) UNIV ILLINOIS FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Streptomyces atroolivaceus.
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                                         06-DEC-2001
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3.0%; Score 12.8; DB 1; Length 18; 87.5%; Pred. No. 3.6e+02; rative 0; Mismatches 2; Indels

2 reacceceácearec 17

(first entry)

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The present invention relates to the isolation of the Streptomyces arroolivaceus leinamycin (Lnm) biosynthesis gene cluster containing 71 controlivaceus leinamycin (Lnm) biosynthesis gene cluster containing 71 can dolored frames (ORFS) (ORFS) - 35 through -1.0 ORFS inmathrough Inma, and one are adding frames (ORFS) (ORFS) - 15 through leinamout antibiotic containing a molecule and stant-incobial activity against Gram-positive and Gram-mogative bacteria, but not against fungi. The polypeptides encoded by the Lnm biosynthesis content of the bost cell is a bacterium or enkaryotic cell, including a condogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite produced by the host cell. The molecule is an endogenous metabolite or an amino acid, and the polypeptide is a peptide synthetase or amino transferame. The polypeptides encoded by the Lnm gene cluster modules and/or catalytic domains are useful for converting an apo-carriet protein to a holo-carrier or protein. Lnm shows potent antitumour acitivity in tumour models in vivo. The Lnm gene cluster modules and/or catalytic domains are useful for useful alone, or in combination with other active domains to modify various target substrates. The Lnm gene cluster is useful to upregulate condogenous modifical Lnm. Lnm, its analogue, or other polyketide, cendogenous modified Lnm. Lnm, its analogue, or other polyketide, cendogenous modified Lnm. Lnm, its analogue, or other polyketide, content an unwher of disorders depending upon the type of metabolites. Abx34311 represent PCR primers used to amplify individual ORFs of the S. atroolivaceus leinamycin blosynthesis
                                                                                                                 Novel gene cluster responsible for synthesis of leinamycin in
Streptomyces atroolivaceus useful for making various peptide and/or
polyketide, and/or hybrid polypeptide/polyketide metabolites.
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                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 28; 185pp; English.
WPI; 2003-018907/01.
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Med Apr 21 12:35:21 2004
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The present invention relates to the isolation of the Streptomyces atroolivaceus leinamycin (Lnm) biosynthesis gene cluster containing 71 open reading frames (OKFS) (OKFS
                                                                                                                                                                                                                                                                                                                   Novel gene cluster responsible for synthesis of leinamycin in
Streptomyces atroolivaceus useful for making various peptide and/or
polyketide, and/or hybrid polypeptide/polyketide metabolites.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 29; 185pp; English.
22-MAR-2002; 2002WO-US008937.
                                                                                                              (REGC ) UNIV CALIFORNIA.
(KYOW ) KYOWA HAKKO KOGYO KK.
                                                          26-MAR-2001; 2001US-0278935P
                                                                                                                                                                                                 Tang G;
                                                                                                                                                                                                                                                           WPI; 2003-018907/01.
                                                                                                                                                                                                    Cheng Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene cluster
                                                                                                                                                                                                    Shen B,
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; 0 Query Match
3.0%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other; Matches

278 GGGCGCCACCAAGCTG 293 GAGCGCCCAAGCTG 18

δ 셤 0×6×6×6×6×6×6×8×8

ABZ68636 standard; DNA; 18 BP.

ABZ68636;

16-MAY-2003 (first entry)

Primer for extension of K121 antibody heavy chain variable region.

K121 antibody; K121-like antibody; kappa-type myeloma cell; kappa-type multiple myeloma; haematopoietic cell transplantation; apoptosis; kappa myeloma antigen; PCR; primer; ss.

Mus musculus.

WO2003004056-A1

16-JAN-2003 

06-JUL-2001; 2001AU-00006179.

05-JUL-2002; 2002WO-AU000896

(PACM-) PACMAB PIY

Dunn RD, Raison RL,

WPI; 2003-210317/20.

Treating kappa-type multiple myeloma in a subject by administering a K121 -like antibody not conjugated to a toxin or a cytolytic agent.

Example 8; Fig 9d; 65pp; English.

antibody heavy chain variable region. The primers were used to construct antibody heavy chain variable region. The primers were used to construct at X121-like antibody by oligonucleotide assembly using PCR. The K121-like antibody competes with K121 for binding to kappa-type myeloma cells. The K121-like antibody is used in the method of the invention. The specification describes a method for treating kappa-type multiple myeloma in a subject, comprising administering a K121-like antibody which is not conjugated to a toxin or a cytolytic agent. The method is useful for treating kappa-type multiple myeloma, autologous haematopoietic cell transplantation, killing kappa-type myeloma cells in a mixed population of cells and inducing apoptosis in kappa myeloma antigen (KMA) bearing cells

Sequence 18 BP; 4 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

3.0%; Score 12.8; DB 1; Length 18; ilarity 87.5%; Pred. No. 3.6e+02; Conservative 0; Mismatches 2; Indels Local Similarity les 14; Conservat Query Match Best Loca Matches

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RESULT 499

ADD24785 standard; DNA; 18 BP. ADD24785; ADD24785 

(first entry)

15-JAN-2004

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Gaps

Human CYP2D6 mutants G1661C and 1707delT probe H212.

diagnostic; pharmaceutical tolerance; side effect; drug; human;
allelic variability; polymorphism; phase I; phase II;
detoxification mechanism; PCR; primer; probe; NAT2; CYP2D6; CYP1A2;
CYP3A4; mEH; TPMT; MTHFR; paraoxonase; CYP2C9; CYP2C19; CYP2C1; DPD; ss.

Homo sapiens

WO2003018837-AZ.

06-MAR-2003.

22-AUG-2002; 2002WO-EP009386.

24-AUG-2001; 2001DE-01040651 30-APR-2002; 2002DE-01019373

(ADNA-) ADNAGEN AG.

Lustig M; Waschuetza S, Schnakenberg E,

WPI; 2003-290079/28

rage 417

Diagnostic kit, useful for assessing a subject's tolerance of drugs, comprises reagents for determining alleles of genes encoding detoxification enzymes.

Claim 6; Page 17; 156pp; German

This invention describes a novel diagnostic kit for determining tolerance of pharmaceuticals in humans by determining allelic variability of at cleast two polymorphisms of a human enzyme involved in phase I and/or II of the detoxification mechanism in a blood, tissue or other human sample, where tolerance is determined from presence or absence of alleles. The kit comprises two pairs of oligonucleotide primers, in which each pair ampolifies, by PCR, part of a gene for a human detoxification mechanism associated enzyme. The kit may also contain two further pairs of oligonucleotides, serving as probes for detection of amplified DNA segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with segments, especially which generate a different signal in the hybridized and non-hybridized condition. The enzymes detected include NAT2, CYP2D6, CYP1A2, CYP3A4, mEH, TPMT, MTHFR, paraoxonase, CYP2C9, CYP2C19, CYP2E1 or DPD. The kit is used to determine an individual's tolerance of a particular drug, to establish a suitable dose and/or to predict if a subject will show side-effects to a drug. The kit provides minimally invasive, safe or and reliable determination of the metabolic capacity of phase I and/or II enzymes at the molecular level. This sequence represents a probe used in the kit of the kit of the invention.

Seguence 18 BP; 4 A; 3 C; 9 G; 2 T; 0 U; 0 Other;

Gaps ; 0 Query Match
3.0%; Score 12.8; DB 1; Length 18;
Best Local Similarity 87.5%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

15 CTGCGGGTGACCGAGG 30 CAGTGGGTGACCGAGG 17

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RESULT 500 ADE15233/c ID ADE15233 standard; DNA; 18 BP. ADE15233; 

(first entry) 29-JAN-2004 Beer spoilage-associated primer SEQ ID 428.

ss; primer; detection; beer-spoilage; lactic acid bacteria; Gram-negative bacteria; spoilage bacteria.

Megasphaera cerevisiae

WO2002103043-A2

27-DEC-2002.

.9-JUN-2002; 2002WO-EP006808

19-JUN-2001; 2001DE-01029410.

(VERM-) VERMICON AG.

Snaidr J; Beimfohr C,

WPI; 2003-175243/17.

New oligonuclectides, useful for rapid detection of beer-spoilage bacteria by in situ hybridization, are specific for type, genus or species

Claim 1; SEQ ID NO 428; 88pp; German.

This invention describes novel oligonucleotides used in a method for detecting beer-spoilage bacteria in a sample. The bacteria detected include lactic acid bacteria of the genera Lactobacillus or Pediococcus, especially the species L. coryniformis, L. perolens, L. buchneri, L. plantarum, L. fructivorans, L. lindneri, L. dasei, L. brevis or P. damnosus or Gram-negative bacteria of the genera Pectinaus and Generalister, specifically P. frisingensis, P. cerevisiphilus and M. cerevisiae. The oligonucleotides of the invention provide rapid detection of spoilage bacteria (typically within 48 hours, compared with 7-12 days for conventional culture methods), can detect all relevant bacteria in parallel, can differentiate between species of the same genus, and are method of the invention. 

Sequence 18 BP; 3 A; 7 C; 3 G; 5 T; 0 U; 0 Other;

Gaps ö y Match 3.0%; Score 12.8; DB 1; Length 18; Local Similarity 87.5%; Pred. No. 3.6e+02; are 14; Conservative 0; Mismatches 2; Indels Query Match

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ò g RESULT 501 AAA27228

AAA27228 standard; DNA; 19

AAA27228;

20-SEP-2000 (first entry)

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Forward PCR primer for FGF8.

Parkinson's disease; neurodegenerative disorder; PCR primer; FGF8; fibroblast growth factor 8; ss.

Rattus sp.

WO200029550-A2. 25-MAY-2000. 

99WO-US027613. 18-NOV-1999;

98US-00195569. 99US-00425462. 18-NOV-1998; 22-OCT-1999; (CALY ) CALIFORNIA INST OF TECHNOLOGY.

Studer L; Ceste M, Doyle J, Wold BJ, Mckay R,

WPI; 2000-387772/33.

Low oxygen culturing of central nervous system progenitor cells useful in treatment of neurodegenerative disorders.

Example 1; Page 36; 80pp; English.

A method for increasing the differentiation of undifferentiated central nervous system (CNS) cells in culture. This novel method involves culturing the cells in low ambient oxygen conditions. Differentiated CNS cells can be used to treat neurodegenerative diseases such as Parkinson's disease. In order to determine the differentiated phenotype messenger RNA levels can be measured using reverse transcription PCR. This involves using PCR primers specific to certain genes. The present sequence is the forward PCR primer used to monitor the message level of FGF8

Sequence 19 BP; 5 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

06-DEC-2000

AAA72197;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is a PCR primer for the fibroblast growth factor 8 gene (RGP8). It was used in reverse transcription PCR to determine betyression patterns of the RGF8 gene in cultured calls. These cells had been grown in low oxygen conditions, and had differentiated to form various types of neuronal cell. The different expression patterns can be used to determine which set of conditions promotes the differentiation of each type of neurone. The different cell types can be used for tissue transplantation, to treat disorders such as stroke, brain and spinal cord neurodegenerative disorders, Huntington's disease, other neurological disorders and psychiatric disorders parkinson's disease, neurological
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                                                                                                                                                                                                                                                                                                                           Rat; cell differentiation; neurodegenerative disorder; stroke; brain injury; spinal cord injury; Alzheimer's disease; epilepsy; Huntington's disease; Parkinson's disease; neurological disorder; cell transplantation; FGF8; fibroblast growth factor 8; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Culturing of neural crest stem cells useful for treatment of neurodegenerative disorders comprises culturing in low ambient oxygen
                                  Gaps
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3.0%; Score 12.8; DB 1; Length 19; 87.5%; Pred. No. 4e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ceste M, Doyle J, Wold BJ, Morrison SJ, Anderson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 5 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                          Fibroblast growth factor 8 mRNA PCR primer #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    294 GTGAAGGACCTGAGCC 309
                                                                     294 GTGAAGGACCTGAGCC 309
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                                                                                                     drahaddadcchahadcc 19
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AAA72197
ID AAA72197 standard; DNA; 19
                                                                                                                                                                                           AAA30349 standard; DNA; 19
                                                                                                                                                                                                                                                               14-SEP-2000 (first entry)
Query Match
Best Local Similarity 87.5
Matches 14, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus sp.
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The invention relates to a retinoid X receptor-gamma (RXR-gamma) knockout mouse whose germ and somatic cells contain an insertion of an exogenous mouse whose germ and somatic cells contain an insertion of an exogenous content of the RXR-gamma gene (exons 3 and 4) which canced the entire DNA binding domain of RXR-gamma. The invention cell-gamma gene, and also encompasses mammalian particularly calcive RXR-gamma and also encompasses mammalian particularly murine, cell lines which are homozygous or heterozygous for a RXR-gamma gene containing an exogenous DNA insert within exoms 3 and 4. The gene containing an exogenous DNA insert within exoms 3 and 4. The capanists or antagonists using the transgenic mouse or mammalian cell ine. The genetically enginesed mouse and cell line are useful in class of receptors. The mouse and cell line allow the investigation at class of receptors. The mouse and cell line allow the investigation at coth the cellular and in vivo levels of a system that lacks one or more specific isoforms of KXR-gamma. This capability will allow the cellular confined importance of each of the RXR-gamma and its isoforms of retinoic acid-mediated gene expression and its isoforms of retinoic acid-mediated gene expression and tissue specific represent mouse RXR-gamma reverse transgenic mice of the invention. The correspondence is an RT-PCR primer for exon E5 of the mouse RXR-gamma reverse for exon E5 of the mouse RXR-gamma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New genetically engineered mice containing alterations in the gene encoding retinoid X receptor, useful for identifying agonists and antagonists of the receptors and in studying retinoic acid mediated gene
                                                                                                                             Mouse retinoid X receptor-gamma gene; RXR-gamma; exon E5;
DNA binding domain; murine; transgenic animal; RXR-gamma knockout mouse;
drug screening; reverse transcription-PCR; RT-PCR primer; 88.
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3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                        Mouse retinoid X receptor-gamma gene exon B5 RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                     (INRM ) INST NAT SANTE & RECH MEDICALE. (CNRS ) CENT NAT RECH SCI. (UYPA-) UNIV PASTEUR LOUIS. (BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Col 12; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             113 CCGCAGCAAGTACGGC 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 CCACAGCAAGTTCGGC 19
                                                                                                                                                                                                                                                                                                                                           97US-00914256.
                                                                                                                                                                                                                                                                                                                                                                                    96US-0024175P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chambon P, Kastner P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-531490/48.
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                                                                                                                                                                                                                                                               US6093873-A.
                                                                                                                                                                                                                                                                                                      25-JUL-2000.
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AAD19298;

RESULT 504

AAD19298/

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Page 244

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Edinger S, Macdougall JR, Millet I, Ellerman K, Stone DJ; Gerlach V, Grosse WM, Alsobrook JP, Lepley DM, Rieger D, Burgess CE; Casman SJ, Spytek KA, Boldog FL, Li L, Padigaru M, Mishra V; Patturajan M, Shenoy S, Rastelli L, Tchernev VT, Vernet CAM; Zerhusen BD, Malyankar UM, Guo X, Miller CE, Gangolli EA;
hypertensive; haemostatic; cardiant; antianginal; dermatological; immunosuppressive; antinflammatory; virucide; antibacterial; anti-HIV; antiparasitic; antiallergic; antiaschmatic; antirheumatic; antiarthritic; vinorerary; anorectic; antidiabetic; immunomodulator; antipsoriatic; nephrotropic; kerolytic; antidiabetic; cerebroprotective; antibactic; antidiateri; carebroprotective; antideorulsant; antiinfertility; antimanic; antidepressant; metabolic; cytostatic; tranquilizer; analgesic; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides the protein and coding sequences of several novel human proteins, designated NOVX. These can be used in the treatment of, amongst others, cancers, autoimmune diseases, infections, inflammatory diseases, storage disorders, muscle disorders, neurodegenerative diseases and developmental defects. The present sequence is a PCR primer or probe used to isolate the sequences of the invention. All of the probes are modified at the 5' end by TBT and at the 3' end by TBMRA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated polypeptide, designated NOVX, useful for treating or preventing in NOVX-associated disorders e.g. cardiomyopathy, atherosclerosis, diabetes, cancer, allergy, asthma, Crohn's disease.
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3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-NOV-2000; 2000US-0253834P.
30-NOV-2000; 2000US-0250926P.
25-JAN-2201; 2001US-0264180P.
20-AUG-2001; 2001US-0313656P.
05-OCT-2001; 2001US-0327456P.
28-NOV-2001; 2001US-00327456P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-590741/63.
                                                                                                                                                                                                                                                                                                                                                                                    WO200257450-A2.
                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The patent discloses a method of screening mammals for autoimmune discases by identifying polymorphisms in interleukin (II)-12 pd0 gene. The methods and kits of the invention are used for screening individuals, families and populations for disease conditions or predispositions for the development of a disease condition which is characterised the development of a disease condition which is characterised are used to treat, prevent or diagnose autoimmune diseases such as IDDM (Insulin dependant diabetes mellitus). The present DNA sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; NOVX; autoimmune disease; cancer; infection; inflammatory disease; storage disorder; muscle disorder; neurodegenerative disorder; nootropic; developmental defect; neuroprotective; antiparkinsonian; hypotensive;
                                                                                                                                                                                                                                                                                                                                                                                Interleukin-12; IL-12 p40; autoimmune disease; Th1/Th2 dysregulation; therapy; allelic variant; insulin dependant diabetes mellitus; IDDM; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening mammals for autoimmune diseases such as diabetes, comprises identifying polymorphisms in interleukin (IL)-12 p40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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3.0%; Score 12.8; DB 1; Length 19;
Best Local Similarity 87.5%; Pred. No. 4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human NOVX coding sequence PCR primer SEQ ID NO: 131.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 5 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                  Mammalian IL-12 p40 intron 7 allelic variant #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
replace(10, A)
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 21; Page 42; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      118 GCAAGTACGGCATGCT 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-MAR-2000; 2000AU-00006466
15-MAY-2000; 2000US-0204366P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-MAR-2001; 2001WO-AU000340
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                                                                         1298/c
AAD19298 standard; DNA; 19
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04-OCT-2001

Mammalia Key allele Morahan G;

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24-OCT-2002

ABT06307/
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AC ABT0
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Huma
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KW Huma
KW Stork
KW deve

ABT06307;

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Wed Apr 21 12:58:21 2004
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WO200183715-A2.
 Homo sapiens
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(LUME/) I
(STUD/) §
(MCKA/) N
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01-MAY-2001; 2001WO-US014051 01-MAY-2000; 2000US-0201005P US GOVERNMENT. LEE S. LUMELSKY N. STUDER L. MCKAY R D G. WPI; 2002-049345/06.

Studer L, Mckay RDG Lumelsky N,

Culturing cells such as neuronal cells for use in treating neurological disorders, comprises generating embryoid bodies from undifferentiated embryonic stem cells, selecting precursor cells, expanding and differentiating them.

Example 10; Page 41; 66pp; English.

The invention provides a method of culturing cells. The method involves expanding a culture of undifferentiated embryonic stem (ES) cells, ceparating embryoid bodies (EB), culturing the bodies to select for central nervous system (CMS) precured cells (PC), culturing PC in an expansion medium comprising a neurologic factor, and differentiated neuronal cells. The method is useful for culturing undifferentiated neuronal cells. The method is useful for culturing undifferentiated neuronal cells which are useful for treating and centrophic factor (BDNF), bFGF, glial derived growth factor (GDNF) and NT-4/5 can be introduced into a brain of a gene product such as tyrosine hydroxylase, nerve growth factor (GDNF), brain derived neurotrophic factor (BDNF), bFGF, glial derived growth cator (GDNF) and NT-4/5 can be introduced into a brain of a subject. The method is useful for culturing dopaminergic, cholinergic and servicenergic neuronal cells. The differentiated neuronal cells are useful continuing epilepsy, familial dysauchosis, severe selzure disorders including epilepsy, familial dysauchosis, severe selzure disorders including epilepsy, familial dysauchosis, severe selzure disorders continuing drug acuse capilla of regulating from aging. Assays are useful for developing drugs capable of regulating from aging. Assays are useful for dopamine or serotonin. Cell cultures comprise 20-40% dopaminergic neurons and 1-3% astrocytes are useful for equences Alforders recomprise and release, particularly for serotonin and opamine, neuronal cells survival, and the electrophysicohemical properties of differentiated neuronal cells. Comprise 20-40% dopaminergic specific regulatory genes used for examining the mechanism of ES cells

Sequence 19 BP; 5 A; 5 C; 7 G; 2 T; 0 U; 0 Other;

; 0 Query Match 3.0%; Score 12.8; DB 1; Length 19; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels

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RESULT 507 ABS97846 ID ABS9

ABS97846 standard; DNA; 19 BP.

(first entry) 23-DEC-2002

Human sulfotransferase thermolabile (STM) gene PCR primer #1.

WO200257410-A2.

25-JUL-2002.

28-NOV-2001; 2001WO-US044838.

28-NOV-2000; 2000US-00724389.

DNAS-) DNA SCI LAB INC.

Guida M, Hall J;

WPI; 2002-698522/75.

Isolated nucleic acid molecules having polymorphisms in known human genes eg. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related tratts.

Example 17; Page 131; 714pp; English.

This invention relates to the sequence of an isolated nucleic acid
molecule comprising at least one base variation from that of a known
that of world the page of (CYP4501A1), cytochrome P450 2 (CYP4501A2),
cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1),
cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1),
cytochrome P450 02E1 (CYP45002E1), adrenergic receptor betal (ADBR1),
cytochrome P450 02E1 (CYP45002E1), proceed transferase (ADBR1), explorated by droxylase 2 (EPHX2), 5-11poxyganase activating inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-11poxyganase activating transferase (NMMT), (AAIlktein 2) KLK2, nicotinanide -N-methyl
cransferase (NMMT), (AAIlktein 2) KLK2, nicotinanide -N-methyl
cransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4
(UGT2B14), UDP-glucuronosyl transferase 2B7
(UGT2B15), urokinase receptor (UPA), multidrug resistance 1
(MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3
(MRR1), orphan nuclear receptor (NR12), or acetylcholine muscarinic
receptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4 or CHRR8) sequence
creeptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4 or CHRR8) sequence
creeptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4 or CHRR8) sequence
creeptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4 or CHRR8) sequence
creeptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4 or CHRR8) sequence
creeptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4, or CHRR8) sequence
creeptor 1, 2, 3, 4, or 5 (CHRR1, CHRR2, CHRR3, CHRR4, CHRR3) contained to crait molecules comparising the genes that
care responsible for specific traits within the genome and eventually
creeptor and/or treating the disorders. The nucleic acid molecules comparising the
collymorphic sequences contained in CYP4501A1, CYP4501A2,
creeptor and/or MDR3 are useful for screening individuals for altered drug

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metabolism. The polymorphic sequences contained in CYP4501A1, CYPP4501A2, AHR, WDR1 and/or MDR3 may also be used to screen individuals for susceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are used to screen for altered cardiovascular function, in COX2 for altered susceptibility to colorectal tumours, in DB1 or CHMR1 for altered central nervous system function, in PLAP and HMM7 for altered pulmonary, immunological or haematological function, in LTF for altered serine procease activity in the prostate, in LTF for altered immunological or haematological function, in CHMR3, CHMR4 for altered central peripheral nervous system function. The present sequence represents a PCR primer used to amplify the sequences of the invention
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Sequence 19 BP; 4 A; 3 C; 7 G; 5 T; 0 U; 0 Other;

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3.0%; Score 12.8; DB 1; Length 19;
87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indels 0; Gaps
                                      14; Conservative
                      Similarity
Query Match
Best Local 8
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55 CAGAGGAGTCTCTGCA 70 2 cacaccacrircrica 17 ð

ABZ69849 standard; RNA; 19 BP (revised)
(first entry) 27-OCT-2003 10-APR-2003 ABZ69849; ABZ69849/C
1D ABZ69849/C
1D ABZ698-C
XX ABZ69-C
XX ABZ698-C
XX ABZ

Ribozyme; Rz2; pharmaceutical carrier; haematopoietic; anti-HIV; virucide; cytostatic; antianaemic; cardiant; gene therapy; cell therapy; antiannes therapy; HV; haemoglobinopathy; leukocyte; Fanconi's anaemia; chronic granulomatous disease; Gaucher's disease; GGFD deficiency; cardiovascular disease; HIV-1-HXB2; ss.

HIV-1 strain HXB2 RNAi target sequence 2.

Human immunodeficiency virus 1.

WO2003006691-A1.

23-JAN-2003

10-JUL-2002; 2002WO-US021907.

10-JUL-2001; 2001US-0304127P. 21-DEC-2001; 2001US-0343484P.

(GENE-) GENE SHEARS PTY LTD.

Macpherson J, Fanning G, Gerlach W; Sun L, Amado R, WPI; 2003-221763/21. Symonds GP,

New composition comprising CD34 hematopoietic cells transduced with a viral construct expressing an anti-HIV agent, useful for treating HIV, AIDS, and diseases of the blood and immune systems, e.g. Fanconi's anemia or cancer

Example 5; Page 112; 157pp; English.

The invention relates to a novel composition comprising a pharmaceutical carrier and haematopoietric cells transduced with a viral construct expressing an anti-HIV agent. A composition of the invention has virucide, cytostatic, antianaemic, anti-HIV, and cardiant activity. The compositions may have a use in gene therapy, cell therapy, and antisense therapy. The composition is useful in the manufacture of a medicament for the treating HIV. The composition can also be used in the treatment of a

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variety of diseases in which there is a genetic aspect, such as diseases of the blood and immune systems, including haemoglobinopathies, defects of leukocyte production or function including cancers, AIDS/HIV, viral infections, lysosomal storage diseases and stem cell defects such as Fanconi's anaemia, chronic granulomatous diseases, Gaucher's disease, GGFD deficiency, and cardiovascular diseases. The present sequence represents a highly conserved RNAi target sequence from HIV-1 HXB2. (Updated on 27-OCT-2003 to standardise OS field)
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Seguence 19 BP; 5 A; 6 C; 5 G; 3 T; 0 U; 0 Other;

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0
3.0%; Score 12.8; DB 1; Length 19; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels
                                       14; Conservative
                   Local Similarity
   Query Match
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ACF35801,

ACF35801 standard; DNA; 19 BP.

06-NOV-2003 (first entry)

Human GPR43 receptor DNA amplifying gene-specific forward primer.

GPR43; G-protein coupled receptor; fatty acid; antiemetic; antimigraine; neuroleptic; antidepressant; tranquilliser; neuroprotective; noctropic; antiparkinsonian; antientimentic; analgesic; cytostatic; metabolic; immunomodulator; antiasthmatic; cardiant; hypotensive; osteopathic; antianginal; antiulcer; antiallergic; cerebroprotective; human; RT-PCR; primer; ss.

Homo sapiens.

WO2003057730-A1.

06-JAN-2003; 2003WO-EP000042,

07-JAN-2002; 2002US-0346396P 

(EURO-) EUROSCREEN SA.

Le Poul E, Detheux M, Brezillon S, Lannoy V, Parmentier M;

WPI; 2003-598359/56.

Identifying agent that modulates GPR43 function, useful for treating migratine, soltzophrenia, anxiety, by measuring binding of GPR43 polypeptide to short chain fatty acid in presence and absence of candidate modulator.

Example 2; Page 80; 136pp; English.

The invention relates to identifying an agent that modulates function of d-protein coupled receptor GPRA3. The method involves measuring the binding of GPRA3 polypeptide to short chain fatty acid [II] in presence and absence of [III], or measuring signaling activity of SPRA3 contacted with [II] in presence and absence of [III], or measuring signaling activity of GPRA3 in presence of [II] and comparing the activity to activity measured in a sample in which GPRA3 is contacted with [II] in presence of [II] and comparing the activity of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating polency of GPRA3 in a call and for modulating will also of disorders such as womiting, migraine, schizophrenia, manic depression, anxiety, dementia, neurodegenerative diseases such as

Alzheimer's disease and Parkinson's diseases and dyskinesias, such as thutlington's disease. They are also useful for preventing, improving or correcting dysfunction or disease s.g., pain, cancer, anorexia, bulimia, asthma, acute heart failure, hypertension, osteoporosis, urinary tetention, angina pectoris, myocardial infarction, ulcers, allergies, stroke, and schizophrenia. The present sequence represents a GPR43 genespecific primer used in semi-quantitative RT-PCR reactions

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Sequence 19 BP; 2 A; 7 C; 5 G; 5 T; 0 U; 0 Other;

0; Gaps Query Match 3.0%; Score 12.8; DB 1; Length 19; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels

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210 RESULT

ADE65585 standard; RNA; 19 BP. ADE65585 ID ADE6

ADE65585; 

29-JAN-2004 (first entry)

Human c-fos transcript target sequence/siNA upper strand, SEQ ID NO:40.

RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairbin RNA; siRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; central nervous system disorder; Alzheimer's disease; parkinson; disease; muntington's disease; pringepsy; dementia; amyotrophic lateral sclerosis; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; transplant rejection; viral infection; HIV infection; autoimmune disease; transplant rejection; vasortopic; nootropic; antiparkinsonian; neuroprotective; cytostatic; antiinflammatory; antialergic; virucide; anti-HIV; immunosuppressive; anticonvulsant; nephrotropic; human; c-fos; target sequence; ss.

Homo sapiens.

WO2003070914-A2.

28-AUG-2003.

20-FEB-2003; 2003WO-US005162.

20-FEB-2002; 2002US-0358580P. 11.MAR-2002; 2002US-0363124P. 06-7UN-2002; 2002US-0366782P. 29-AUG-2002; 2002US-0406378P. 05-SEP-2002; 2002US-0408378P.

(SIRN-) SIRNA THERAPEUTICS INC.

09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P.

Beigelman L; Mcswiggen J,

WPI; 2003-679877/64.

New short interfering nucleic acid downregulates expression of the c-fos gene useful for treatment and diagnosis of diseases, e.g. cancer and inflammation.

Example 3; SEQ ID NO 40; 145pp; English.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human c-fos gene by RNA interference. The

cc stranded. They further comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or clearatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA), the siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the c-fos gene in cells, tissue explants or organisms of siNA, by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating central nervous system lesions and injuries (e.g., Alzheimer's disease, explants or alternative diseases, epilepsy, dementia or amyotrophic lateral sclerosis and polycystic kidney disease); inflammatory conditions and allergic diseases; virtal infections (including HIV infection); autoimmune diseases; virtal infections (modulance autoimmune diseases; and transplant rejection. The siNAs are also useful function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human c-fostarded double-stranded siNA, which is identical to the c-fos transcript target sequence.

%XGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG

Sequence 19 BP; 6 A; 5 C; 6 G; 0 T; 2 U; 0 Other;

. 0 3.0%; Score 12.8; DB 1; Length 19; 75.0%; Pred. No. 4e+02; ative 2; Mismatches 2; Indels Local Similarity 75.0 les 12; Conservative Query Match Best Local Si Matches 12;

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286 CCAAGCTGGTGAAGGA 301 

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ADE65701 standard; RNA; 19 RESULT 511 ADE65701/c

BP.

29-JAN-2004 (first entry) ADE65701;

Human c-fos siNA lower strand, SEQ ID NO:156.

RNA interference; short interfering nucleic acid; siNA; shNA; shNA; shNA; shNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; central nervous system disorder; Alzheimer's disease; Parkinson; disease; huntington's disease; pilepsy; dementia; amyotrophic lateral sclerosis; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; transplant rejection; viral infection; HIV infection; autoimmune disease; transplant rejection; vasorropic; antiparkinsonian; neuroprotective; cytostatic; antiinflammatory; antiallergic; virucide; anti-HIV; immunosuppressive; anticonvulsant; nephrotropic; human; c-fos; ss. 

Homo sapiens.

WO2003070914-A2.

28-AUG-2003.

10-FEB-2003; 2003WO-US005162

20-FEB-2002; 2002US-0358580P. 11-MAR-2002; 2002US-0363124P. 06-UJN-2002; 2002US-0386782P. 29-AUG-2002; 2002US-0406784P.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human c-fos gene by RNA interference. The siNAS may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense coligonucleotide. Specifically, the siNAS include short interfering RNA (siRNA). The siNAS can be unmodified or chemically modified, can contain ceoxyribonucleotide, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits of the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAS are used to modulate expression of the c-fos gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating central nervous system lesions and injuries (e.g., Alzheimer's disease, Parkinson's disease, Huntington's disease, epilepsy, dementia or amyotrophic lateral sclerosis, various cancers; other proliferative ciseances; and polycystic kidney disease); inflammatory and/or allergic diseases; wiral infections (including HIV inflection); autoimmume diseases; and transplant rejection. The sinAs are also useful function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human c-fos-New short interfering nucleic acid downregulates expression of the c-fos gene useful for treatment and diagnosis of diseases, e.g. cancer and Example 3; SEQ ID NO 156; 145pp; English. (SIRN-) SIRNA THERAPEUTICS INC. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. WPI; 2003-679877/64. Mcswiggen J, inflammation. 05-SEP-2002; 

Sequence 19 BP; 2 A; 6 C; 5 G; 0 T; 6 U; 0 Other;

Gaps .. Match 3.0%; Score 12.8; DB 1; Length 19; Local Similarity 87.5%; Pred. No. 4e+02; les 14; Conservative 0; Mismatches 2; Indels Query Match Best Loca Matches

CCAAGCTGGTGAAGGA 301

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ABZ86355 standard; DNA; 20 BP ABZ86355; RESULT 512 ABZ863 

Human oligonucleotide sequence (first entry)

17-0CT-2003

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; ofvostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

31-OCT-2002.

24-APR-2001; 2001US-0286137P.

Beigelman L;

WPI; 2003-229219/22.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oilgonucleotide antisense to the intration codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an artifulflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a municense gene therapy. The composition is useful for treating or preventing a respiratory, lung activity. The composition may have a composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an extinctlammatory steroid in a subject, for reducing levels of adenosine or receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp. WIPO.

Sequence 20 BP; 5 A; 6 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.8; DB 1; Length 20;
Best Local Similarity 87.5%; Pred. No. 4.5e+02;
Matches 14; Conservative 0; Mismatches 2; Indels

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16 Argreadchdercegr 1 ઠે g AAT06919 standard; DNA; 19 BP. AAT06919; BXBXSXXXXXXXXXXXXXXXXX

Chromosomal locus E17 primer #1.

Synthetic.

WO9532214-A1

30-NOV-1995

WO200285308-A2

WEG APT 41 14.30.41 400

23-APR-2002; 2002WO-US013135. 

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; (EPIG-) EPIGENESIS PHARM INC. Nyce JW, I Miller S,

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

claim 15; SEQ ID NO 1597; 872pp; English.

190 ATATCCACTGCTCGGT 205

RESULT 513 AAT06919 04-JUL-1996 (first entry)

prostate/colon tumour suppressor gene; allelic loss; colorectal cancers; microsatellite analysis; sequence tagged site; primer; probe; PCR; amplification; chromosome; ss.

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95WO-US006593

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WPI; 1997-108751/10.
                                WPI; 1996-020526/02.
                                                                                                       Sest Local Similarity
                 (CANJ-) CANJI
                         Bookstein R,
                                                                                                                                                                                             WO9702048-A1
                                                                                                                                                                                                            28-JUN-1996;
                                                                                                                                                                                                                              24-AUG-1995;
28-SEP-1995;
09-APR-1996;
                                                                                                                                                           19-OCT-1997
   22-MAY-1995;
          20-MAY-1994;
                                                                                                                                                                                                                                          12-APR-1996;
                                                                                                                                                                                                     23-JAN-1997
                                                                                                                                                                                                                   30-JUN-1995
20-JUL-1995
                                                                                                                                                                                                                           26-JUL-1995
                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                         Kleyn PW,
                                                                                                                                                    AAT48575;
                                                                                                   Query Match
                                        New DNA
                                                                                                           Matches
                                                                                                                                     RESULT 51
AAT48575/
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Nucleic acid sequence determination - comprising synthesising chain extension products, which are indicative of positions of selected species of nucleotide in nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a primer for exon 3 of the HLA-C gene. This sequence can be used in the method of the invention for determining the position of at least one selected species of nuclectide, in a region of interest, in a sample. The method comprises combining the sample with a reaction mixture to synthesise chain extension products indicative of the positions of the species of nuclectide in the region of interest and evaluating the products produced, characterised in that the sample, which is combined with the reaction mixture, and contains target and non-target nucleic acid polymers in natural abundance. The method can be used to detect mutations, particularly mutations of medical significance, in samples
                                                           The murine and human tub gene (AAT48550 and AAT48551 respectively) products are wild-type, expressed in the hypothalamus. The form lacking exon 5 is produced by alternative splicing. The products participate in the control of mammalian body weight. Measuring tub expression and detection of tub gene mutation are used to diagnose body weight disorders, esp. obesity, cachexia and anorexia, or related sensory and fertility defects
                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detection;
human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shipman
                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                      3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; amplify; pathogen identification; mutation uncleic acid analysis; microorganism characterisation; HLA type determination; HLA-C gene exon 3; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lacroix J,
                                                                                                                                                                                                                                  Seguence 19 BP; 6 A; 6 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5' vgicwsp5 primer for exon 3 of HLA-C gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Larson MT,
                                                                                                                                                                                                                                                                                                               0; Mismatches
                                  Disclosure; Page 35; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 24; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                        132 CIGGCCCGCCTGGCGGTGG 150
                                                                                                                                                                                                                                                                                                                                                                                              19 chracchachachad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96US-00640672.
96US-00684498.
97US-00807138.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97WO-US007135.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT99886 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                              Local Similarity 78.5
les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-549755/50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-1996;
19-JUL-1996;
27-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leushner J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9741259-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT99886;
                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                    Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT
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                                                                                                                                                                                                                                                                                                                 Primers AAT06887-932 were used to analyse the breakpoints at chromosomal locus 892-21, contained in patients having prostate cancer, by microsatellite analysis and sequence taggaed sites (STS). The region contains a prostate/colon tunour suppressor gene (PTS). The primers and amplified fragments were used to screen a YAC library of prostate cancer DNA to isolate the PSTG (AAT06880), which can be used in the diagnosis and treatment of prostate and colorectal cancers. The primers AAT06919-20 amplify a 121 bp fragment from chromosomal locus E17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid encoding mammalian tub protein - useful for diagnosis and treatment of body wt. disorders, esp. obesity, and for screening for
                                                                                                                                                                                                                   , DNA encoding a prostate tumour suppressor protein - from chromosome for the diagnosis and treatment of prostatic and colorectal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tubby; tub; CBT9 gene; body weight; obesity; cachexia; anorexia; disorders; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 7 A; 6 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                 Disclosure; Page 86; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            182 CAAGGCACATATCCACTGC 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAAGGCATATCACAACTGC 19
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95US-0001444P.
95US-0002759P.
95US-0004424P.
96US-0015396P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96WO-US011186
                                                             94US-00246604
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human tub gene primer R12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15; Conservative
                                                                                                                                          Isaacs WB;
                                                                                                    INC.
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The present primer was used in the mapping of a gene encoding 2 forms of a prostate/colon tumour suppressor (P/CTS). The P/CTS gene was isolated by analysis of allelic loss in patients with prostate cancer, and was putatively located to the chromosomal location 8g22-21 via microsatellite analysis and the use of sequence tagged sites (STS). Primers and probes derived from the gene can be used to screen lambda cDNA libraries for genes encoding P/CTS form 1 and 2. The P/CTS or its cDNA can be used in the diagnosis and treatment of prostate and colorectal cancers. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct
                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prostate/colon tumour suppressor; allelic loss; prostate cancer; colorectal cancer; microsatellite analysis; sequence tagged site; STS; amplification; chromosomal location 8q22-21; probe; primer; gene mapping;
derived from a human patient, animal, plant or microorganism, determine HLA type ancillary to transplant procedures, detect and identify microorganisms, particularly pathogenic microorganisms, in a sample and in in situ sequencing reactions to produce sequencing fragments in a histological specimen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New prostate/colon tumour suppressor gene - mapped to a locus on human
                                                                                                                                                                     Gaps
                                                                                                                                                                     ò
                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer E17 for mapping prostate/colon tumour suppressor gene.
                                                                                                                               3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 7 A; 6 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 26; 45pp; Japanese.
                                                                                                                                                                                                     378 GACCGCGACGACGCCCCA 396
                                                                                                                                                                                                                           GACCGCGGGGGCGGGCCA 19
                                                                                                  Sequence 19 BP; 2 A; 6 C; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96JP-00062144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-00445515
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(UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                        AAT64713 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                     Local Similarity 78.9
nes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosis; treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isaacs WB, Bookstein R;
                                                                                                                                                                                                                                                                                                                                                                                             (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chromosome 8
                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
12-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP09098790-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                           AAT64713;
                                                                                                                                                                                                                                      -
                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                          RESULT 516
                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                            AAT64713
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This sequence represents a PCR primer for the human ACE (angiotensin converting enzyme) gene, and can be used in the method of the invention.

The method is for assessing cardiovascular status in humans by the determining the sequence of at least one polymorphic site in the ACE (angiotensing enzyme), ACT (angiotensing and/or ATI (type 1 angiotensin converting enzyme), ACT (angiotensing and/or ATI (type 1 converting enzyme), ACT (angiotensing and the polymorphic pattern angiotensin II receptor) genes, and comparing the polymorphic pattern of the tin patients with predetermined markers of status. The method is used to assess blood pressure or electrocardiographic profile, to diagnose a cardiac condition such as (silent) myocardial infarction (MI), the prestrension, atherosciarcosis or stroke. They can also be used to predict response to treatments with ACE inhibitors, angiotensin II receptor antagonists, diuretics, alpha- or beta-adrenergic receptor antagonists, etc. It is also used to identify susceptibility to cardiovascular conditions in the genes are used to screen for cardiovascular agents. The nucleic acids containing to the peptides from the genes are used to screen for cardiovascular agents. The nucleic acids contained in the library can be is used as source of probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Assessing cardiovascular status in humans by polymorphic analysis - of genes for angiotensin converting enzyme, angiotensinogen and angiotensin II receptor, used to diagnose predisposition to disease and to predict effect of therapy.
                                                                                                                                                                                                                                                                                 PCR primer; human; ACB; angiotensin converting enzyme; angiotensinogen; cardiovascular status; AGT; ATI; type I angiotensin II receptor; stroke; polymorphic pattern; blood presenue; electrocardiographic profile; cardiac condition diagnosis; mycoardial infarction; atherosclerosis; hypertension; cardiovascular disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gabs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 7 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Norberg LT, Andersson MK, Lindstroem PHR;
                                                                                                                                                                                                                                                       Primer ACE/82RB for human ACE gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 27; 71pp; English.
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182 CAAGGCACATATCCACTGC 200
                                    1 caaddcarárcacaacide 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      darricricacciccide 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98WO-IB000475.
                                                                                                                                       AAV08577 standard; DNA; 19
                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EURO-) EURONA MEDICAL AB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-568361/48.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9845477-A2
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                                                                                                                                                                                                                   15-FEB-1999
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Best Local Si
Matches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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                                                                                                                                                                           AAV08577;
                                                                                               RESULT
AAV0857
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Gaps

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Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels

WO9920749-A1.

Synthetic

AAX16754;

RESULT 518

AAX16754,

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The specification describes a method for screening for functional polypeptides which bind a ligand. The method comprises contacting a repertoire of polypeptides with a generic ligand, and then screening selected functional polypeptides with a target ligand. The method permits the removal from a chosen repertoire of polypeptides, those which are non-functional, e.g. as a result of the introduction of frame-shift mutations, stop codons, folding mutants or expression mutants which would be or are incapable of binding to any target ligand. The method also permits the enrichment of a chosen repertoire of polypeptides for those polypeptides which are functional, well foldied and highly expressed. The polypeptides obtained can be used in diagnostic, prophylactic and therapeutic procedures. PCR primers AAX35946-48 were used to amplify a general in which was used in the construction of libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                          Screening for functional polypeptides which bind a ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 2 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer for PG1 biallelic marker 99-123-184.
                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 49; 67pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              270 CIGGAGCAGGGGGCACCA 288
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98US-0099658P.
                                                                         98WO-GB003135
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97US-0066729P.
                                                                                                                                                                                           (MEDI-) MEDICAL RES COUNCIL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15; Conservative
                                                                                                                                                                                                                                     Tomlingon I, Winter G;
                                                                                                                                                                                                                                                                        WPI; 1999-288302/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PSA; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-SEP-1998;
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                                                                                                                                                       21-NOV-1997;
                                                                           20-OCT-1998;
                                                                                                                 20-OCT-1997;
                                   29-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ01311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primers AXX16733-X16754 are examples of primers which can be used to PCR amplify the human "tub" gene (AAX16702) exons. The invention relates to a method for identifying compounds that modulate tub protein activity, especially its interaction with proteins containing an SN2 domain. The method can be used for identifying compounds which modulate tub protein activity for use in the treatment of mammalian body weight disorders including obesity, cachexia and anorexia
                                                                                                                                                                                                                  Mouse, wild type, tubby, identification, SH2 domain, mammal, obesity, body weight disorder, cachexia, anorexia, primer, PCR, amplification, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying compounds which modulate tub protein activity - by detecting compounds which alter the interaction of tub protein with a SH2 containing peptide, used to develop agents for treating e.g. obesity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Screening, functional polypeptide; ligand, non-functional, enrichment, single chain antibody, PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5' primer used to amplify germline V gene segment DP-47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 6 A; 6 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                              Human tub gene exon 12 R12 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Col 22; 95pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kleyn PW;
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97US-00829553.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX35946 standard; DNA; 19
                                                            AAX16754 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                          27-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kapeller R, Moore KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cachexia or anorexia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-130383/11.
                                                                                                                                                                                                                                                                                                    Homo sapiens.
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AAX35946;

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RESULT

AAX35946,

Query Match

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Gaps ö rng.res

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The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polymucleoride and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allele specific amplification assay for determining the identity of a nucleotide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate specific antigen (BSA) diagnosis of prostate cancer relies on prostate specific antigen (BSA) and postate specific antigen (BSA) inability to discriminate between malignant and non-malignant affections of the organ. A need exists for both a reliable diagnostic procedure curative treatments of the disease. The PG1 gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Claim 4; Page 367; 385pp; English.
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Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                       289 AGCTGGTGAAGGACCTGAG 307
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Human HPC2 cDNA exon 24 3'UTR mutation screening primer SEQ ID NO: 185 AAA60364 standard; DNA; 19 (first entry) 07-DEC-2000 AAA60364; RESULT 52 

Human, mouse; prostate cancer predisposing gene; HPC2; human chromosome 17p; gene therapy; peptide therapy; drug design; PCR primer; sequencing primer; ss.

40200027864-A1. domo sapiens

99WO-US026055 05-NOV-1999; 18-MAY-2000

98US-0107468P. 06-NOV-1998;

(MYRI-) MYRIAD GENETICS INC.

Teng DHF, Simard J, Rommens JM; WPI; 2000-376481/32. Tavtigian SV,

Human prostate cancer (HPC)2 nucleic acids, polypeptides, and antibodies, useful for treatment and diagnosis of prostate cancer.

Example 5; Page 62; 157pp; English

The present sequence is a primer used in the isolation of the human and murine prostate cancer predisposing genes HPC2 and Mm.HPC2. The human

ö version of the gene is found on chromosome 17p. Some alleles cause a predisposition to cancer, particularly prostate cancer. This gene and its protein can be used in peptide and gene therapy for cancer patients, as wall as being useful as diagnostic tools (both for cancer sufferents and those with a predisposition to the disease) and in the production of cancer drugs Gaps ö Length 19; 4; Indels G; 0 T; 0 U; 0 Other; Query Match
3.0%; Score 12.6; DB 1;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; CCACACAGAGGAGCCACAG 19 CCACTCAGAGGAGTCTCTG 68 Sequence 19 BP; 7 A; 7 C; 5 20

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Bougueleret L;

Chumakov I,

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Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss. cdk4 ribozyme binding site #10. 묡. AAA82829 standard; DNA; 19 (first entry) WO200032765-A2. 04-DEC-2000 Mammalia. AAA82829; RESULT 522 AAA82829/c

cleaves New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1. Robbins JM; Barber JR, 99WO-US028772. 98US-0110954P. Tritz R, Welch PJ, WPI; 2000-412314/35. (IMMI) IMMISOL INC. 06-DEC-1999; 04-DEC-1998; 08-JUN-2000 

Disclosure; Page 52; 109pp; English.

The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in Gaps ; 0 Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels Sequence 19 BP; 1 A; 7 C; 5 G; 6 T; 0 U; 0 Other; restenosis treatment

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75 GAGGGCGCGCAGTGGACA 93 19 gagggccacaaagrggcca 1 ઠે

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523 RESULT

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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDKI, PCNA and Cyclin BI. Representative examples of ribozyme recognition sites are given in AAAB2415 to AAAB46787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment
                                                                                                                  Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                  New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Angiotensin-converting enzyme gene, ACE, polymorphism, polymorphic marker, cardiovascular disease, myocardial infarction, unstable angina; hypertension, atherosclerosis, stroke, prognosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human angiotensin-converting enzyme (ACE) PCR primer, SEQ ID NO:2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                screening; treatment outcome; human; PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 3 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                           Tritz R, Welch PJ, Barber JR, Robbins JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 85; 109pp; English.
                                                                                    Cyclin F ribozyme binding site #221.
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AAA38202 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-AUG-2000 (first entry)
AAA84953 standard; DNA; 19
                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 78.9
les 15; Conservative
                                                                                                                                                                                                                                                                                               (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-412314/35.
                                                                                                                                                                              WO200032765-A2.
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                                                          04-DEC-2000
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                                                                                                                                                                                                                                                                    04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA38202;
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                             AAA84953;
                                                                                                                                                Mammalia.
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AAA38202/c
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The invention relates to a novel method of assessing the cardiovascular cratus in an individual and to newly identified polymorphisms in the eratus in an individual and to newly identified polymorphisms in the common and construction of an angiotensian. Converting enzyme (ACE), angiotensian II receptor type 1 (ATI) and type 2 (ATI), angiotensiangen (ACI), renin, aldosterone synthase, endothelin receptor type A and beta-adreneragic receptors; and 2. The method comprises determining the sequence at one or more polymorphic positions within these genes, and comparing the crospication appointable from the individual with a reference polymorphic pattern obtained from a population of individuals exhibiting a pattern obtained from a population of individual to cardiovascular disease status. The polymorphic markers are predetermined cardiovascular status of a patient given a predicting the likely cardiovascular status of a patient given a treatment regimen comprising administration of cardiovascular drugs (e.g., ACE inhibitors, beta-adreneracic receptor antagonists (beta-conflorate and probes for detecting penetic polymorphic site may be used as the provides a basis for predicting the outcome of atreatment regimen. Tedimes of the genes comprising a polymorphic site may be used as primares and probes for detecting genetic polymorphisms or in molecular contrast arrays for high throughput soriening. The genes, and the proteins of they encode are useful in the screening of potential cardiovascular dungs a petermination of an individual soriening. The spenses of aliminate patient from the intail with a pub-population of potential can be evaluated to cardiovascular patients from collinical trials who are predicted to a particular cegimen Adverse results in an early trial can be evaluated with a sub-population of the test population, permitting correlated with a sub-population of the treatment group. Beneficial correlated with a sub-population and evers response, to a particular correlated with a sub-population and ever of sup
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 7 A; 3 C; 7 G; 2 T; 0 U; 0 Other;
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AAA09605/c
ID AAA09605 standard; DNA; 19 BP.
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AC AAA09605;
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Best Local Similarity 78.9
Matches 15, Conservative
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Assessing cardiovascular status in humans involves comparing test polymorphic pattern comprising polymorphic positions within genes encoding specific proteins, with reference polymorphic pattern.

Example 1; Page 48; 126pp; English

Jonsson

Norberg LT, Andersson MK, Lindstrom PHR,

WPI; 2000-318010/27

99WO-IB001678 98US-0104286P.

AAA84953/c

14-OCT-1998; 14-OCT-1998; 13-OCT-1999;

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Novel phosphodiesterase and its gene for research on complex mechanism of intracellular information transfer.
                                                                                                                                                                                                                                                                                                            Sequences AAA09589-A009592 encode human phosphodiesterase 10 (PDE10) proteins AAB26853-B26856. Phosphodiesterase 10 and its gene are useful for research on the complex mechanism of intracellular information transfer. The invention includes a recombinant vector containing a PDE10 gene, and a cell transfermed with the vector. Sequences AAA09593-A09606 represent PCR primers used in the isolation of the PDE10 polynucleotide sequences of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ACE, AGT and AT1 genes polymorphisms PCR primer SEQ ID NO: 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism, disease diagnosis, treatment; cancer; system; nervous system; glaucoma; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                        3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                Phosphodiesterase 10; PDE10; human; PCR primer; ss
                        PCR primer SEQ ID 17 used in PDE10 identification.
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 2 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                         Example 6; Page 27; 29pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGTGCACCTGGAGCAGGGC 281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGTCCACCTGGAAGAGCGC 1
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99WO-IB000497.
99US-0126243P.
99US-00471890.
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                                                                                                                                                                             98JP-00338861.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
 (first entry)
                                                                                                                                                                                                   (TANA ) TANABE SEIYAKU CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
hes 15; Conserva
                                                                                                                                                                                                                          WPI; 2000-605129/58.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; genetic
cardiovascular
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                                                                                                   JP2000224992-A.
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24-MAR-1999;
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                                                                                                                            15-AUG-2000.
                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               263
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Query Match
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

360 GACTTCCTCACTTTCCTGG 378

GATTICITCACCICCTGG 1

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AAC71201 standard; DNA; 19

RESULT 527

AAC71201;

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Gaps

; 0

Sequence 19 BP; 7 A; 3 C; 7 G; 2 T; 0 U; 0 Other;

methods of the invention

The present invention is related to methods for determining the polymorphic pattern of an individual and using the results to determine their risk of a number of diseases, including cancer, cardiovascular diseases, glaucoma and nervous system disorders such as depression and neurodegenerative diseases. In addition, the methods can be used to determine the effects of different types of treatment for individuals, and thus enables appropriate therapies to be prescribed. THE PCR primers shown in sequences AAC61201-C61371 were all used to demonstrate the

Assessing disease status in individual by determining sequence(s) at one or more polymorphic positions within the human genes encoding the protein(s) involved in physiological pathway associated with treatment

WPI; 2000-638268/61.

Example 1; Page 55; 141pp; English.

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Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                     Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human
of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in genes. These SNPs can be used in disease diagnosis and prediction
                                                                                                                                                                                                                                                                                                                                   Lander ES;
                                                                                                                                                                                                                                                                                                                                , Daley GQ, Ireland JS,
Sklar P;
                                                               Single nucleotide polymorphism PCR primer #688.
                                                                                                                                                                                                                                                                                          (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 8; Fig 5; 214pp; English.
                                                                                                                                                                                                                                       30-MAR-2000; 2000WO-US008440.
                                                                                                                                                                                                                                                                  99US-0127248P.
                                       (first entry)
                                                                                                                                                                                                                                                                                                                                     Altshuler D, Cargill M,
Lipshutz RJ, Patil N, 9
                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-611722/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    genetic analysis.
                                                                                                                                                                                 WO200058519-A2
                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                    31-MAR-1999;
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                                       09-FEB-2001
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Sanders

Olaisson E,

117 AGCAAGTACGGCATGCTGG 135

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individual's susceptibility to disease, in forensic and paternity testin and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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Patil N, Sklar P;
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llarity 78.9%; Pred. No. 4.4e+02;
Conservative 0; Mismatches 4; Indels
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3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                                                                                       Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism PCR primer #720.
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                                                                                                                                                                                                                                                   117 AGCAAGTACGGCATGCTGG 135
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Matches 15; Conservat
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The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose usefulbility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, echizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
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Lipshutz RJ, Patil N, Sklar P,
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                                                                                                                                                                                  Single nucleotide polymorphism PCR primer #666.
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1 AGCACGTGAGGCATTCTGG
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                                                                                      AAC71168 standard; DNA; 19
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Best Local Similarity 78.9
Matches 15; Conservative
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                                           Single nucleotide polymorphism; SNP; human; genetic disease;
disease susceptibility; cardiovascular system; endocrine system;
neurological system; forensic testing; paternity testing; PCR primer; ss.
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disease susceptibility, cardiovascular system; endocrine system,
neurological system; forensic testing; paternity testing; PCR primer; ss.
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Lipshutz RJ, Patil N, Sklar P;
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Single nucleotide polymorphism PCR primer #724.
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Best Local Similarity 78.9
Matches 15, Conservative
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The present invention is concerned with a number of human single mucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
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                                                                                               Cargill M, Daley GQ, Ireland JS, Lander ES; Patil N, Sklar P;
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Sklar P;
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(WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
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Patil N, S
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Matches 15; Conserv
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Lipshutz RJ,
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genetic analysis
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                                                                             Homo sapiens
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Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other; AAC71198; diseases RESULT 534 AAC71198 ሯ 쉱 ប្ដង្គន ò 셤 The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose usceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's ö The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, echizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. Single nuclectide polymorphism, SNP; human, genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss. for phenotypic correlations, forensics, paternity testing, medicine and Gaps ö Altshuler D, Cargill M, Daley GQ, Ireland JS, Lander ES; Lipshutz RJ, Patil N, Sklar P; 3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels Sequence 19 BP; 4 A; 4 C; 7 G; 4 T; 0 U; 0 Other; Single nuclectide polymorphism PCR primer #700. (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC. 117 AGCAAGTACGGCATGCTGG 135 AGCACGTGAGGCATTCTGG 19 Claim 8; Fig 5; 214pp; English Claim 8; Fig 5; 214pp; English BB 30-MAR-2000; 2000WO-US008440. 99US-0127248P AAC71219 standard; DNA; 19 (first entry) 15; Conservative WPI; 2000-611722/58. Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in diesase diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, schizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system; neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                   Gaps
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Patil N, Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels
         Length 19;
                                                                4; Indels
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Score 12.6; DB 1;
Pred. No. 4.4e+02;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism PCR primer #686.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           117 AGCAAGTACGGCATGCTGG 135
                                                                                                                                 117 AGCAAGTACGGCATGCTGG 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AGCACGTGAGGCATTCTGG 19
                                                                                                                                                                                           1 Accaccicacccarrered 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 8; Fig 5; 214pp; English.
                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAR-2000; 2000WO-US008440.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0127248P.
         Query Match
Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                             AAC71198 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200058519-A2
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Lipshutz RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-1999;
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rng.res

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Human; WHL gene; sequencing; mutation; human leukocyte antigen; HLA;
transplantation surgery; detection; identification; primer;
pathogenic microorganism; ss.
                                                                                                                                                                                                                                                                                      Bi-directional sequencing of nucleic acid polymers for identifying pathogens or detecting mutations by using a single reaction mixture having first and second primers with different, spectroscopically-distinguishable labels.
                                                                                                                                                                                                                                                     Dunn JM, Larson MT, Lacroix J, Shipman R, Leushner J;
                                                                  Human leukocyte antigen C exon 3 sequencing primer SEQ ID NO:13
                                                                                                                                                                                           96US-00640672.
96US-00684498.
97US-00807138.
97WO-US007134.
                 AAA65792 standard; DNA; 19 BP.
                                                                                                                                                                           98US-00009483
                                                                                                                                                                                                                                       (VISI-) VISIBLE GENETICS INC
                                                   (first entry)
                                                                                                                                                                                                                                                                        WPI; 2000-464336/40.
                                                                                                                                                                                             01-MAY-1996;
19-JUL-1996;
27-FEB-1997;
29-APR-1997;
                                                                                                                         Homo sapiens
                                                                                                                                                                            10-JAN-1998;
                                                   22-NOV-2000
                                                                                                                                         JS6083699-A.
                                                                                                                                                           04-JUL-2000
                                  AAA65792;
                                                                                                                                                                                                                                                         Hui M,
535
RESULT 53
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The present invention describes a method for simultaneously determining the position of a nucleotide base in a target region of both strands of a denatured duplex nucleic acid polymer. The method comprises using a single set of reaction mixture that is combined with the nucleic acid polymer. The reaction mixture contains first and second oligonucleotide primers, each with different, spectroscopically-distinguishable contained in samples derived from a human patient, medically significant mutations, in samples derived from a human patient, animal, plant or microorganism, and for the determination of human contains (the proposed of the method can also be used to detect and identify microorganisms, especially pathogenic microorganisms, in a sample, and in situ sequencing pathogenic microorganisms, in a sample, and in in situ sequencing can cancino so produce sequencing fragments within a histological specimen, which are then removed from a selected location on the tissue preparation and loaded onto a gel for sequence analysis. The sequencing reaction is contoured of a disease condition is known, but the causative mutation is contour. The present sequence represents a sequencing primer for the human that C gene, which is used in an example from the present invention Example 2; Col 11; 27pp; English.

Sequence 19 BP; 2 A; 6 C; 11 G; 0 T; 0 U; 0 Other;

Gaps ; 0 3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels Matches 15; Conservative Query Match Best Local Similarity

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à

RESULT 536 AAC85428

AAC85428 standard; cDNA; 19 BP.

AAC85428;

(first entry) 20-APR-2001 Primer oML69 amplifies Salmon MHC class II beta and alpha promoters

Promoter; regulation; expression; MHC class II; RAN; U2A'; primer; immune response; productivity; DNA vaccination; cytokine; amplify; interferon gamma; beta-carotene; polymerase chain reaction; PCR; ss

Synthetic.

WO200077232-A1.

21-DEC-2000.

09-JUN-2000; 2000WO-NO000202.

10-JUN-1999; 99NO-00002819

(GENO-) GENOMAR AS.

Syed M, Lundin M;

WPI; 2001-080695/09.

Novel promoters from Atlantic salmon for regulating expression of nucleotide constructs, as DNA vaccines for protecting salmon and other fish species against viral, bacterial infections.

Example 1; Page 7; 29pp; English.

The sequences given in AACS\$425-28 are primers which were used in the complification of the Salmon WHC class II beer and alpha promoters and introduction of the Salmon WHC class II beer and alpha promoters and introduction of the amplified sequences may be incorporated in nucleotide constructs for the purpose of regulating the expression of such constructs. For example, the salmon WHC class II beta-promoter was a nestred in a plasmid vector carrying the lac2 gene. The resulting constructs for promote transcription of the Lac2 gene. The resulting plasmid vactor carrying the had gene of the resulting plasmid vactor and plasmid vactor almoners almon presembles. The remults showed that fishes injected with the WHC class II promoter results showed that fishes injected with the WHC class II promoter results showed that fishes injected with the WHC class II promoter. Containing plasmid, showed an ELISA titer higher than the mean value +2 times the standard deviation of the control fishes. No immune response was detected using DNA-plasmid with Lac2 gene without promoters. Constructs such as these, may be used in vivo to achieve productivity enhancement in production organisms which are bony fish aquatic and constructs may also be useful for DNA vaccination of salmon and other constructs may also be useful for DNA vaccination of salmon and other fishers and may also be combined with WHC alleles for optimizing the expression and presentation of pathogenic application of pathogenic and pression and presentation of pathogenic and control of salmon are specific outsides and may also be combined with WHC alleles for coloring in salmonids. The promoters in salmon are specific and coloring in salmonids. The promoters is active. The constitutive coloring the expression and presentation of the gene of interest constitutive of promoters halps in expressing the gene of feter-acoteness and compounds responsible for restricted to cells in where expression of the gene of interest constitutively in all colors to the viral infection si 

Sequence 19 BP; 4 A; 4 C; 6 G; 5 T; 0 U; 0 Other;

Gaps ö Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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EP1067191-A2.

.0-JAN-2001.

05-APR-2001

AAF31028;

09-JUL-1999;

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The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme (I) which cleaves RNA encoding a cytckine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antiposciatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, dynamous or basal cell carichmam and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, virreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal deteachment, and for treating and preventing proliferation and second as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57877 to AAH62099 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; appid dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar; sickle cell retinopathy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 3.0%; Score 12.6; DB 1; Length 19; Local Similarity 78.9%; Pred. No. 4.4e+02; les 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 1 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 102; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75 GAGGGCGCGCAGTGGACA 93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH60115 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Robbins JM, Tritz R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-300427/31.
                                                                                                                                                                                                                                                               WO200130362-A2.
                                                                                                                                                                                                                                                                                                                                                                                                        26-OCT-1999;
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                                                                                                                                                                                            sapiens
                                                                                                                                                                                                                                                                                                              03-MAY-2001:
                                                                                                                                                                                                                  Synthetic.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is a PCR primer for wild-type leuk gene from E.coli, which encodes alpha-isopropylmalate synthase (IPMS). The leuk gene was used to generate a mutant alpha-IPMS, which is de-sensitised in feedback inhibition by L-leucine. The mutant alpha-IPMS is useful for the production of L-leucine, which is useful for medical treatment, as a production of other amino acids such as Jysine. The present sequence was used to amplify the leuk gene for use in the present invention
                                                                                                                                                                                                                                                                                                                           Alpha-isopropylmalate synthase; enzyme; IPMS; leucine production; leuA; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cell-cycle dependent kinase cdk4 ribozyme binding site SEQ ID NO:415.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptide with alpha-isopropylmalate synthase activity and decreased feedback inhibition of activity by L-leucine, useful for production of L-leucine for medical treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lunts MG, Kozlov YI, Ivanovskaya LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 19 BP; 2 A; 4 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 9; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  275 GCAGGGGGCACCAAGCTG 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCACATCGCCACCAAGCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99RU-00114325.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUL-2000; 2000EP-00114458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH57991 Standard; DNA; 19
                                                                                                                                               AAF31028 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                        leuA gene PCR primer LeuA9
                                                                                                                                                                                                                                         (first entry)
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Best Local Similarity 78.9'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AJIN ) AJINOMOTO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-125730/14.
                                                                                                                                                                                                                                                                                                                                                                                                        Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gusyatiner MM,
Voroshilova EB;
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Gaps

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Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;

Cyclin F ribozyme binding site SEQ ID NO:2539.

Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy;

10-SEP-2001

RESULT 538
AAH57991/C
ID AAH57991/C
XX
AC AAH5
DT 10-S
DX
XX
XX
XX
Huma
KW reco

AAH57991;

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538

Homo sapiens. Synthetic.

26-0CT-1999;

03-MAY-2001

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The present invention relates to a number of prostate-specific sequences derived from the human PS118 gene. These can be used in the detection, monitoring and treatment of prostate diseases, particularly prostate cancer. The PS118 fragments of the invention were isolated from a prostate tissue expressed sequence tag (EST) library. The present sequence is a PCR primer used to isolate a sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel PS118 polypeptide for detecting, diagnosing, staging, monitoring, prognosticating, preventing, treating, or determining predisposition of individual to diseases and conditions of prostate, e.g. prostrate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, mouse, HPC2, prostate cancer, neoplastic growth, cytostatic, si
gene therapy, prostate cancer predisposing gene, chimpanzee, gorilla,
sequencing primer, PCR primer.
                       Human, prostate, prostate-specific sequence, prostate cancer, PS118; cytostatic, gene therapy, PCR; primer, 8s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.0%; Score 12.6; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 4.4e+02; Matches 15; Conservative 0; Mismatches 4; Indels
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Klass MR, Kratochvil JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 3 A; 2 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 41; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          139 GCCTGGCGGTGGAGGCCGG 157
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                                                                                                                                                                                                                                               26-NOV-2001; 2001US-00991681.
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98US-00065383.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Billingel PA, Cohen M,
Granados EN, Hodges SC,
Russell JC, Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                                  (COLP) COLPITES T L. (FRIE) FRIEDMAN P N. (GORD) GORDAN J. (GRAN) GRANADOS E N. (KLAS) KLASS M R. (KRAY) KRATOCHVIL J D (KORY) KORSCHIL J C. (KUSS) RUSSELL J C. (KTRO) STROUPE S D. (STRO)
                                                                                                                                                                                                                                                                                                                                                                      BILLINGEL P A.
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                                                                                                                                                    US2002086316-A1.
                                                                                                                                                                                                                                                                                                                                                                                                COHEN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
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Russell JC,
                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                23-APR-1997;
23-APR-1998;
                                                                                                                                                                                                  04-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                    (BILL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HOMO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method for treating a proliferative skin or eye disease and scarring. The method involves administering a ribozyme [1] which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (WMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule [1] comprising a promoter operably linked to a nucleic acid segment encoding [3]. [4] can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. [7] can be used in gene therapy. [1] and [1] are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH52737 to AAH62099 represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammation, matrix metalloproteinases, growth factors and cell-cycle dependent kinases.
recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipisoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 19 BP; 3 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 256; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           321 GIGCIGGCGGCGACGACC 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 GTGCTGACGCAGGAGTACC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL49572 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                             6-OCT-2000; 2000WO-US029500.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Robbins JM, Tritz R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IMMU-) IMMUSOL INC.
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Gordan J; Roberts-Rapp L;

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Gaps

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AAL49572 RESULT 540
AAL49572
ID AAL4957
AC AAL4957
XX
DT 27-NOVDT 27-NOVDE Human g

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(FARB ) BAYER AG

Smolyar A;

Rommens JM;

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G-protein coupled receptor; receptor; serotonin; 5-hydroxytryptamine; human; antibacterial; virucide; fundicide; protezoacide; neuroprotective; cardiant; antidepressant; hypertensive; hypotensive; diuretic; costeopathic; antiulcer; antihalammatory; antiallergic; cytostatic; nootropic; analgesic; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a human prostate cancer predisposing gene coding for an HPC2 polypeptide. The DNA and protein sequences are useful as diagnostic reagents for identifying a mutant HPC2 nucleotide sequence in a suspected mutant HPC2 allele by comparing the sequence of the suspected mutant HPC2 allele with a wild-type HPC2 sequence. The sequences are also useful for detecting an alteration in HPC2, where the alteration is associated with cancer in a human. The method involves analysing an HPC2 gene or an HPC2 gene expression product from a tissue of the human. The HPC2 gene is useful as a marker for prostate cancer and can be used in gene therapy techniques to suppress neoplastic growth of recipient cells which carry the mutant HPC2 allele. The sequences represent primers used in the methods of the invention, CDNA encoding human and mouse HPC2 and
                                                                                                                                                                                                                                                                                     Novel nucleic acid sequence encoding HPC2 polypeptide, which is marker for prostate cancer, is useful in gene therapy techniques to restore HPC2 normal levels by which neoplastic growth is suppressed in recipient cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human serotonin-like G-protein coupled receptor PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 7 A; 7 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in the methods of the invention, cDNA encoding cDNA encoding HPC2 paralogues and orthologues
                                                                                                                                                                                                                 Tavtigian SV, Teng DHF, Simard J,
                                                                                                                                                                                                                                                                                                                                                                 Example 8; Page 75; 239pp; English.
                                                                                                                                                           (MYRI-) MYRIAD GENETICS INC.
(HOSP-) HOSPITAL FOR SICK CHILDREN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCACTCAGAGGAGTCTCTG 68
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2001US-0324054P.
                                                                                       07-MAY-2001; 2001WO-US014602.
                                                                                                                          05-MAY-2000; 2000US-00564805.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABN84896 standard; DNA; 19
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                                                                                                                                                                                                                                                    WPI; 2002-066599/09.
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                   WO200185911-A2
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24-SEP-2001;
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                                                    15-NOV-2001
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ABN84896/c
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                                                                                                                                                                                                                                                                                                                                                                                                                      cancer, a urinary disorder, obesity, diabetes, a central nervous system (CNS) disorder, asthma or a hematological disorder (all claimed) in a subject. The reagent is especially an antisense oligonucleotide, ribozyme or antibody. Pharmaceutical compositions comprising the reagent, or an expression vector encoding 5HT-like GPCR, are claimed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                         New G-protein coupled receptor (GPCR) polynucleotide and its encoded protein, useful for identifying modulators of GPCR activity, and in gene therapy for treating bacterial infection, cancer, acute heart failure or Parkinson's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human chromosome 1p36-35 PCR primer SEQ ID NO:2078.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 2 A; 7 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                    Example 21; Page 124; 164pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         284 CACCAAGCTGGTGAAGGAC 302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ilarity 78.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (RIKA ) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL45034 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-144136/19.
                                                                               WPI; 2002-643344/69
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Matches
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Gaps 0

3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; Live 0; Mismatches 4; Indels

BP

Detecting presence of target PS118 polynucleotide in test sample, useful for detecting, diagnosing, staging, monitoring, prognasticating, preventing or treating or determining predisposition to prostate disease.

WPI; 2002-187683/24.

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The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates numbered for discrimination are mixed from the arker of a sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the maximum in the specified discrimination Nos. to array the multiwell containing the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell containination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the calculate are specified from the amplified by using the above primer; (g) signals resultant cultures are amplified by using the above primer; (g) signals created efform the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The mixed plates are specified from the detected result; and (i) the clones are precipied from the detected result; and (i) the clones are precipied for human chromosome 21q223; which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PS118; prostate; marker; prostate cancer; human; sequencing; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Roberts-Rapp L;
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0
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Klass MR, Kratochvil JD,
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 5 A; 7 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prostate-specific PS118 clone sequencing primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              87 GIGGACATCACCACGICTG 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Grecacarcaccaracere 19
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Hodges SC,
Stroupe SD;
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HODGES S C.
KLASS M R.
KRATOCHVIL J D.
ROBERTS RAPP L.
RUSELL J C.
STROUPE S D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BILLINGEL P A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
nes 15; Conserv
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Granados EN, B
Russell JC, St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GORDON
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                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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(COHE/)
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(KLAS/)
(KRAT/)
(ROBE/)
(RUSS/)
(STRO/)
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(FRIE/)
(GORD/)
(GRAN/)
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X88888888888888888888888888888
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The present sequence is that of a sequencing primer designed from sequencing information of a prostate-specific PS118 consensus sequence tages ABA91561): It was used in the sequencing of PS118 expressed sequence tag-specific clones (see ABA91642-50) transcribed from human prostate tissue. PS118 polypeptides (see AAMS0809-13), polymucleotides (see aAMS0809-13), stagings) monitoring, prognosticating, preventing and treating, or determining the predisposition of an individual to, hyperplasia, prostatic intraepithelial neoplasia, prostate cancer, tumours and metastases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lolium perenne; perennial ryegrass; plant; cell wall; lignification; cellulase; enzyme; lignin biosynthesis; cellulose degradation; CCOAMT; caffeoyl-CoA 3-O-methyltransferase; cinnamyl alcohol dehydrogenase; CAD; caffeic acid O-methyltransferase; OMT; cinnamete-4-hydroxylase; C4H; cinnamoyl-CoA reductase; CCR; percoxidase; PER; ferulate-5-hydroxylase; F5H; CELL; phenylalanine ammonia lyase; PAL; 4-coumarate:COA ligase; 4CL; ryegrass; fescue species; molecular genetic marker; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid encoding lignification and cellulase enzymes or their related enzymes useful for modifying lignin biosynthesis and cellulose degradation in plants to manipulate plant cell wall.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ong EK, Emmerling M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 3 A; 2 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AGRI-) AGRIC VICTORIA SERVICES PTY LTD.
(AGRE-) AGRESEARCH LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lolium perenne LpPeroxidasel primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 37; 436pp; English
                                                                                                                                                                                                    Example 2; Page 41; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     139 GCCTGGCGGTGGAGGCCGG 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cacricicida de caracia de cacricia de cacr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Spangenberg G, Sawbridge TI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN87259 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-SEP-2001; 2001WO-AU001221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-SEP-2000; 2000AU-00000419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-444025/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200226994-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lolium perenne.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-APR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABN87259;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABN87259
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The present invention describes a nucleic acid (I) or its fragment encoding caffeoyl-CoA 3-O-methyltransferase (CCAMT), cinnamyl alcohol dehydrogenase (CAD), caffeid acid 0-methyltransferase (OMT), cinnamate-4-hydroxylase (CRI), peroxidase (PER), cinnamoyl-CoA reductase (FCR), peroxidase (PER), cinnamate-4-lydroxylase (FCR), peroxidase (PER), collulase (CELL), fernlate-5-hydroxylase (FSF), peroxidase (PER), collulase (CELL), fernlate-5-hydroxylase (FSF), peroxidase (PER), collulase (CELL), fernlate-5-hydroxylase (FSF), peroxidase (PER), collulase (PER), collula
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3.0%; Score 12.6; DB 1; Length 19; Similarity 78.9%; Pred. No. 4.4e+02; Conservative 0; Mismatches 4; Indels 392 CGCCAAGAAGGTCTTCTAC 410 1 CGCCAAGAAGAACCTCAAC 19 15; Conservative Query Match Best Local S 셤 ઠ

ABZ76924 standard; DNA; 19 BP 07-MAY-2003 (first entry) ABZ76924; RESULT 546 ABZ76924 

Homo sapien Synthetic.

16-JAN-2003.

05-JUL-2002; 2002WO-EP007520.

06-JUL-2001; 2001EP-00116412. 13-MAY-2002; 2002US-0379412P.

Fries H, Winter A;

WPI; 2003-239205/23.

New nucleic acid molecule comprising a sequence of an allele of a polymorphic bovine acyl CoA-diacylglycerol transferase gene useful for testing a mammal for its predisposition for fat content of milk and for meat marbling.

Example 2; Page 27; 91pp; English.

The present invention describes a nucleic acid molecule (NA) (I) encoding to a bovine acyl CoA-diacylglycerol transferase (DGAT) contributing to or indicative for low fat content of milk and to low meat marbling correction for low fat content of milk and to chromosome 8, and (intramuscular fat content). Human DGAT is located to chromosome 8, and content of milk and/or its and its predisposition for fat content of milk and/or its correction for meat marbling. The method comprises analysing the gene conding DGAT for nucleotide polymorphisms (e.g. single nucleotide correction for meat marbling. The method comprises analysing the gene conding DGAT for nucleotide polymorphisms are located in the coding region of the DGAT conclected polymorphisms are located in the coding region of the DGAT conclected has at the position 10433 and 10434 of the DGAT conclected has at the position 10433 and 10434 of the DGAT conclected has at the position of an amino conclected milk and low meat marbling. The nucleic acid molecule has at the position of the DGAT corresponding to position 10433 and 10434 of the DGAT corresponding to position 10433 and 10434 of the DGAT corresponding to position 10433 and 10434 of the DGAT corresponding to position 10433 and 10434 of the DGAT gene and high meat marbling. The nucleic acid molecule has at the position of regidues which correlate with a predisposition for high content of milk and high meat marbling. The prodisposition for high content of milk and high meat marbling the appreasion of the expression of the present conduct of the gene encoding DGAT. ABZ7045 to ABZ77045 and ABP96035 to high content sequences used in the exemplification of the present invention

Sequence 19 BP; 2 A; 7 C; 5 G; 5 T; 0 U; 0 Other;

0; Gaps Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

> ઠે 원

ACC62358 standard; DNA; 19 BP. ACC62358; RESULT 547 ACC62358

23-JUN-2003 (first entry)

Human NOV5 forward PCR primer SEQ ID NO:233.

Human; NOVX; antiatherosclerotic; hypotensive; cardiant; dermatological; anorectic; immunosuppressive; cytostatic; antidiabetic; antifitaterility; heamostatic; antiinflammatory; antiasthmatic; anti-HIV; immunomodulator; neuroprotective; noctropic; antiparkinsonian; metabolic; antilipaemic; gene therapy; cardiomyopathy; atherosclerosis; hypertension; soleroderma; consential heart defect; aortic stenosis; valve disease; transplantation; tuberous sclerosis; obesity; congenital adrenal hyperplasis; diabetes; prostate cancer; metabolic disorder; neoplasm; lymphoma; uterus cancer; idiopathic thrombocytopenic purpura; AIDS; bronchial asthma; anorexia; crohn; disease; multiple sclerosis; infectious disease; cancer; ancer anc

Homo sapiens. Synthetic

WO2003023001-A2

20-MAR-2003,

09-SEP-2002; 2002WO-US028538.

07-SEP-2001; 2001US-0318120P. 07-SEP-2001; 2001US-0318184P.

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rng.res

Sequence 19 BP; 8 A; 7 C; 3 G; 1 T; 0 U; 0 Other;

Gaps ;

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Human DGAT gene forward PCR primer 1534.

Acyl CoA:diacylglycerol transferase; DGAT; enzyme; chromosome 8; human; milk; meat marbling; low fat; polymorphic; SNP; single nucleotide polymorphism; PCR primer; ss.

sapiens.

WO2003004630-A2.

(ARBE-) ARBEITSGEMEINSCHAFT DEUT RINDERZUECHTER.

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10-SEP-2001; 2001US-0318430P.
17-SEP-2001; 2001US-0322636P.
17-SEP-2001; 2001US-0322816P.
17-SEP-2001; 2001US-0322816P.
19-SEP-2001; 2001US-0322816P.
20-SEP-2001; 2001US-032361P.
20-SEP-2001; 2001US-032363P.
25-SEP-2001; 2001US-032363P.
25-SEP-2001; 2001US-032496P.
25-SEP-2001; 2001US-032496P.
26-SEP-2001; 2001US-032499P.
14-PES-2002; 2001US-0329599P.
16-PES-2002; 2001US-0395999P.
17-WAY-2002; 2002US-0381863P.
25-MAY-2002; 2002US-0381863P.
                                                                                                              06-SEP-2002; 2002US-00236417.
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wed Apr 21 12:38:21 2004

### (CURA-) CURAGEN CORP.

Agee ML, Alsobrook JP, Anderson DW, Berghs C, Boldog FL;
Burgess CE, Casman SJ, Catterton E, Chant JS, Chaudhuri A;
Crabtree J, Dipippo VA, Edinger SR, Eisen AJ, Ellerman K;
Gangolli EA, Gerlach VL, Gotte L, Gorman L, Guo X, Gusev VY, Ji W;
Kekuda R, Khramtsov NV, Leach MD, Lepley DM, Li L, Liu X;
Malyankar UM, Miller CE, Ooi CE, Ort T, Padigaru M, Patturajan M;
Pena CEA, Rieger DK, Rothenberg ME, Shenoy SG, Shimkets RA;
Spaderna SK, Spytek KA, Taupier RJ, Twomlow N, Vernet CAM, Voss EZ;
Zerhusen BD, Zhong M;

#### WPI; 2003-313241/30.

Novel human proteins and nucleic acid encoding the proteins, useful for diagnosis, treatment and prevention of disorders involving the human protein or nucleic acid e.g. cardiac and neurological disorders.

### Example C; Page 301; 460pp; English.

The present invention describes isolated human NOVX proteins, where X is 1 to 42. ACC62236 to ACC62345 encode the human NOVX proteins given in ABR54276. NOVX sequences have antiatheroscelerotic, cardiant, hypotensive, dermatological anosectic, immunosuppressive, cytostatic, antidiabetic, antiinfertility, haemostatic, antiinflammatory, anti-HIV, antisathmatic, metabolic, immunomodulator, neuroprotective, nootropic, antiparkinsonian and antilipaemic activities, and can be used in gene therapy. NOVX proteins are useful for treating or preventing a pathology associated with a NOVX protein in humans and for trating a syndrome associated with the human disease. NOVX nucleic acids, proteins and antibodises can be used in the treatment and diagnosis of cardiomyopathy, atherosclerosis, hypertension, congenital heart defects, actic stenosis, valve disease, tuberous sclerosis, scleroderma, obesity, transplantation, congenital heart defects, actic stenosis, congenital adrenal hyperplasia, prostate cancer, fertility, haemophilia, disconders, neoplasm, lymphoma, uterus cancer, fertility, haemophilia, disease, allos bronchial asthma, Crohm's disease, multiple sclerosis, and mecatoolic disorders, dyslipidaemias, and metabolic syndrome X. Alzheimer's disease, parkingon's disease, multiple sclerosis, and mematopoietic disorders, dyslipidaemias, and metabolic syndrome X. Acc62465 represent PCK primers and probes for human NOVX processent PCA 277 processent processis of conferences, which are used in examples from the present invention. ABK54277 represents a human trypsinogen protein given in comparison with the human NOV35b protein in the exemplification of the present invention

# Sequence 19 BP; 6 A; 3 C; 9 G; 1 T; 0 U; 0 Other;

Gaps ô Query Match
3.0%; Score 12.6; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 4.4e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anrediabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gasrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; psoriasis; inflammatory bowel disease; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss.
                                                                                                               Mitogen activated protein kinase sinA oligonucleotide SEQ ID NO:414
260 caccercaccrccaccac 278
         1 CAGGGAGGACCTGGAGAAG 19
                                                           ADE29792 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                 28-JAN-2003; 2003WO-US002510
                                                                                              29-JAN-2004 (first entry)
                                                                                                                                                                                                                                WO2003072590-A1.
                                                                                                                                                                                                                                                 04-SEP-2003.
                                                                                                                                                                                                               Synthetic.
                                                                             ADE29792;
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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes. WPI; 2003-689980/65.

Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira

(SIRN-) SIRNA THERAPEUTICS INC.

20-FEB-2002; 2002US-0358580P. 11-WAR-2002; 2002US-0365124P. 06-UJN-2002; 2002US-0366784P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 15-JAN-2003; 2003US-0409233P.

## Example 3; SEQ ID NO 414; 164pp; English.

The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a mitogen-activated protein kinase that downregulates expression of a mitogen-activated protein kinase ((NAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or corganisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) conjugates and/or complexes of siNA; and (4) vectors that express siNA and cells containing these vectors. MAPK siNAs containtritic, ancrectic, antibacterial, antirhelumator, antisathmatic, immunosuppressive, antibacterial, antirhelumator, antisathmatic, immunosuppressive, antibacterial, antirhelumator, containts and activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obseity, diabetes types I condition and validation; psociasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target containts, they can also be used for drug screening; diagnosis; target containts, and containts, pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide collantification and gene mapping (e.g. of single-nucleotide in the exemplification of the present invention.

Sequence 19 BP; 4 A; 8 C; 4 G; 0 T; 3 U; 0 Other;

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;
                          Gaps
                          6
3.0%; Score 12.6; DB 1; Length 19; 68.4%; Pred. No. 4.4e+02; iive 2; Mismatches 4; Indels
                             Conservative
                Local Similarity
                             13;
  Query Match
               Best Loca
Matches
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280 GCGCCACCAAGCTGGTGAA 298 GCUGCCCCAACCUGCUGAA 19

Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:510. ADE29888 standard; RNA; 19 (first entry) 29-JAN-2004 ADE29888; ADE29888 

short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; ortookatic; anorecitic; antidabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; psoriasis; inflammatory Dowel disease; drug screening; genetic engineering; paramacogenomic; gene mapping; ss.

Synthetic.

WO2003072590-A1.

04-SEP-2003.

28-JAN-2003; 2003WO-US002510.

11.MAR.2002, 2002US-0363124P. 06-UUN-2002, 2002US-03867B2P. 29-AUG-2002, 2002US-04067B4P. 05-SEP-2002, 2002US-040937BP. 09-SEP-2002, 2002US-0409293P.

(SIRN-) SIRNA THERAPEUTICS INC

15-JAN-2003; 2003US-0440129P

Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B;

WPI; 2003-689980/65.

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated diagnosis of cancer, oprotein kinase genes.

Example 3; SEQ ID NO 510; 164pp; English.

the present invention describes a short interfering nucleic acid (siNh) that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo corganisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) expression of siNA; and cells containing these vectors. MAPK siNAs have cytostatic, anorectic, antidabetic, antifiniammatory, antiarthritic, antipsoriatic and gastrointeserinal activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, eptic shock, rheumatoid arthritis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; pharmacogenomics; cientification and validation; genetic engineering; pharmacogenomics;

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studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK siNA which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                             short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidiabetic; antifiammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; psoriasis; inflammatory bowel disease; drug screening;
                                                                                                                                                                                                                                                                      Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:519
                                                                                               ..
0
                                                                     3.0%; Score 12.6; DB 1; Length 19; 68.4%; Pred. No. 4.4e+02; ative 2; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                  genetic engineering; pharmacogenomic; gene mapping; ss
                                                Sequence 19 BP; 3 A; 6 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                       136 CCCGCCTGGCGGTGGAGGC 154
                                                                                                                                            1 ccueccueaaecuesáese 19
                                                                                                                                                                                                      ADE29897 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                      (first entry)
                                                                                                 13; Conservative
                                                                                     Local Similarity
                                                                                                                                                                                                                                                      29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                              ADE29897;
                                                                          Query Match
                                                                                       Best Loca
Matches
                                                                                                                                                                                 RESULT 550
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Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B; WPI; 2003-689980/65.

(SIRN-) SIRNA THERAPEUTICS INC.

11-WAR-2002; 2002US-0363124P. 06-JUN-2002; 2002US-0366782P. 29-AUG-2002; 2002US-0406784P. 05-SEP-2002; 2002US-0408378P. 09-SEP-2003; 2002US-0408293P. 15-JAN-2003; 2003US-0440129P.

28-JAN-2003; 2003WO-US002510.

WO2003072590-A1.

04-SEP-2003

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes. Example 3; SEQ ID NO 519; 164pp; English.

The present invention describes a short interfering nucleic acid (sinA) that downregulates expression of a mitogen-activated protein kinase (whkpK) genes by RNA interference. Also described: (1) a method for modulating expression of MaPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of sinA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA and cells containing these vectors. MaPK siNAs have cytostatic, anorectic, antidiabetic, antiinflammatory, antiacthritic, antiposriatic and gastrointestinal activities. The MaPK siNAs siNAs can be used to modulate the expression of MAPK genes, in cells,

tissue explants or organisms, e.g. for treating obesity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, septic shock, rheumatoid arthrities, psoriasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target identification and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK sinA which is used in the exemplification of the present invention. 88888888888888

Sequence 19 BP; 3 A; 4 C; 8 G; 0 T; 4 U; 0 Other;

0; Gaps 3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; ive 0; Mismatches 4; Indels 0; GCGCCACCAAGCTGGTGAA 298 15; Conservative Best Local Similarity 280 Query Match Matches ઠ

19 GCTGCCCCAACCTGCTGAA 1 용

ADE29783 standard; RNA; 19 29-JAN-2004 (first entry) ADE29783; RESULT 551 ADE29783/C 

short interfering nucleic acid, siNA, downregulation, inhibition, mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; oytostatic, anridiabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antitheumatic; antiarthritit; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; pesoriasis; inflammatory bowel disease; drug screening; genetic engineering; pharmacogenomic; gene mapping; ss. Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:405.

Synthetic

04-SEP-2003.

WO2003072590-A1.

28-JAN-2003; 2003WO-US002510 20-FEB-2002;

11.-MAR.-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
09-SEP-2002; 2002US-0409378P.
09-SEP-2003; 2003US-0409293P.

(SIRN-) SIRNA THERAPEUTICS INC.

Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B;

WPI; 2003-689980/65.

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes.

Example 3; SEQ ID NO 405; 164pp; English.

The present invention describes a short interfering nucleic acid (siNA) that downregulates expression of a mitogen-activated protein Kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4)

vectors that express sinh and cells containing these vectors. MAPK sinhs have cytostatic, anorectic, antidiabetic, antinflammatory, anorectic, antidiabetic, antinflammatory, antiasthmatic, immunosuppressive, antibacterial, antirhemmatic, antipaemiatic and gastrointestinal activities. The MAPK sinhs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obseity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, septic shock, rheumatoid arthitis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; diagnosis; target definitization and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK sinh which is used in the exemplification of the present invention.

Sequence 19 BP; 3 A; 7 C; 6 G; 0 T; 3 U; 0 Other;

4; Indels 0; Gaps 3.0%; Score 12.6; DB 1; Length 19; 78.9%; Pred. No. 4.4e+02; tive 0; Mismatches 4; Indels Conservative Local Similarity hes 15; Conserv Query Match Best Loca Matches

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136 cccccccccccccccc154 19 CCTGCCTGAAGCTGGAGGC 1 à g

AAZ94278

AAZ94278 standard; DNA; 20 BP.

AAZ94278;

03-JUL-2000 (first entry)

PHBLIX; human; testis-specific; transcription factor; prostate cancer; bladder cancer; ovary cancer; testicular cancer; gene therapy; diagnosis; vaccine; PCR primer; ss. Human PHELIX nested primer NP2.

99WO-US020137. WO200012709-A2. 31-AUG-1999; Homo sapiens. 09-MAR-2000.

98US-0098610P. 98US-0106524P. UROGENESYS INC. AFAR D E. HUBERT R S. , Ю O 31-AUG-1998; 31-OCT-1998; (HUBE/) I (UROG-) 

Raitano AB Afar DE, Hubert RS,

RAITANO A B.

WPI; 2000-237872/20

Testis specific Helix Loop Helix proteins expressed in cancers and useful for the prevention, diagnosis and treatment of prostate, bladder and ovarian tumors.

Example 1; Page 31; 62pp; English

The present sequence is that of nested primer NP2, which was used in the amplification of gene fragments obtained from a suppression subtractive hybridization reaction using LAPC xenograft cDNA and designed to identify novel prostate and prostate cancer-specific genes. A 437 bp clone was obtained. Full-length cDNA (see AAS94275) was subsequently cloned from a transcription factor that is normally expressed only in testis cDNA there is up-regulated in prostate and other types of cancer. The invention

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a PCR primer used in the isolation of cDNA fragments of the PTAN (testis specific protein expressed in prostate cancer) gene. PTAN is expressed in 3 isoforms PTAN-1. 2, and 3. The PTAN gene is located on chromosome 1q22. PTAN is overexpressed in prostate cancer, and has a testis specific expression pattern in adult tissues PTAN shows no homology to any known gene. PTAN can be used in methods for the diagnosis of cancer, especially prostate or breast cancer, where the normal tissue samples are prostate tissue, or breast tissue, bone tissue, lymphatic tissue, serum, blood, or urine. A vector containing the PTAN nucleotide sequence, a vaccine composition targeting PTAN, PTAN treat cancer, especially breast and antisense sequences, can be used to treat cancer, especially breast and prisense sequences, can be used to treat cancer, especially breast and prisense sequences. Cancer development can be inhibited by a vaccine composition targeting PTAN
provides diagnostic and therapeutic methods useful in the management of various cancers which express PHELIX, including prostate cancer, bladder cancer, ovarian cancer and testicular cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PTAN; testis specific; prostate cancer; overexpress; chromosome 1q22; diagnose; cancer; breast; vaccine; PCR primer; ss.
                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PTAN proteins, and sequences encoding them, used for diagnosing and treating cancers, especially breast and prostate cancers.
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                                                                                                     Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                         Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer (NP2) used in PTAN gene isolation.
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                                                                                                                                                                                319 GCGTGCTGGCGGCGGACGA 337
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98US-0102910P.
98US-0113229P.
99US-0129518P.
                                                                                                                                                                                                                                                                                                        AAA37951 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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HUBERT R S.
RAITANO A B.
MITCHELL S C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-317715/27.
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21-DEC-1998;
14-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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(HUBE/)
(RAIT/)
(MITC/)
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                                                                                                                                                                                                                                                                        RESULT 553
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BPC-1 polypeptides and polynucleotides can be used for the detection of BPC-1 polypeptides and polynucleotides in biological samples, this is particularly useful for detecting cancers expressing BPC-1, e.g. procestate cancer or bladder cancer. Antibodies directed against BPC-1 or antisense concer or bladder cancer for treating such cancers. The BPC-1 or antisense polymucleotides can also be used for treating sor inhibiting the careful and predicting susceptibility to developing cancer. The BPC-1 polypeptides and predicting susceptibility to developing cancer. The BPC-1 polypeptide and predicting susceptibility to developing cancer. The BPC-1 polypeptide comprises a cTDB domain which is expressed in prostate and bladder carcinoma cells and which shows sequence similarity with CTB domains from tissues of the brain, however, it is expressed at high levels in certain tissues of the brain, however, it is expressed at high levels in prostate cancer cells and bladder cancer cells. A number of synthetic colligonucleotides were used to generate BPC-1 cDNA from total cell RNA of tumour cells lines. These primars were a cDNA synthesis primer (AAZS3041), two adaptor sequences (AAZS3045), a PCR primer (AAZS3045) and two nested primers (AAZS3047, AAZS3048). This sequence is one of the nested primers (NP)1 used in the amplification method.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated BPC-1 polypeptides, useful for developing products for t
diagnosis, staging, prognosis and treatment of cancers, particularly
prostate or bladder cancer.
                                                                                                                                                                                                                                                                                                             BPC-1; oncogene; oncogenic; cancer; prostate; bladder; antibody; antisense; vaccine; detection; prognosis; drug screening; primer; ss
                    Gape
                                                                                                                                                                                                                                                                             Primer used for generating human brain specific protein BPC-1 cDNA.
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Pred. No. 4.9e+02;
0; Mismatches 4; Indels
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                                                     319 GCGTGCTGGCGGCGGACGA 337
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     Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                              AAZ93048 standard; DNA; 20
                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LEONG K.
RAITANO A B.
SAFFRAN D C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (JAKO/) JAKOBOVITS A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-206006/18.
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HUBERT R S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-AUG-1999;
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                                                                                                                                                                                                                                                 24-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                AAZ93048;
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(AFAR/)
(HUBE/)
(LEON/)
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(SAFF/)
                                                                                                                                               RESULT 554
                                                                                                                                                        AAZ93048
ID AAZ9
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3.0%; Score 12.6; DB 1; Length 20;

Query Match

3.0%; Score 12.6; DB 1; Length 20;

Query Match

**AA**Z94898

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Prostate cancer, testis-specific protein Y-encoded mRNA, TSPY mRNA, vaccine, PCR primer, 8s.
                                                                                                                                                                                                                                         PCR primer for testis-specific protein Y-encoded DNA.
                                    AAA14807 standard; DNA; 20 BP.
                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200020638-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                       08-AUG-2000
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                                                                                                      AAA14807;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 557
AAA14807

AAA148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel testes-specific gene 22P4F11 which is expressed in human prostate cancer and is useful as a diagnostic marker and/or therapeutic target for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22P4F11; human; testis; prostate cancer; diagnosis; gene therapy; marker;
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                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer NP2 used in testis-specific 22P4F11 gene amplification.
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                                    4; Indels
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Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4
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                                                                                                  319 GCGTGCTGGCGGCGGACGA 337
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99US-0146584P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vaccine; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UROG-) UROGENESYS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-303452/26
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28-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-AUG-2000
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PCR primers AAA14805-07 were used to amplify testis-specific protein Yencoded DNA. The specification describes a new method of diagnosis of prostate cancer. The method comprises determining the level of testis-specific protein Y-encoded (1987) mRNA or protein, and comparing these presence of elevated 1987 mRNA or protein is indicative of prostate cancer. Detection of TSPY mRNA expression or protein levels to be comparing the diagnosis of prostate cancer. Antisense polymucleotides complementary to the coding sequence of human TSPY are useful for treating prostate cancer by inhibiting TSPY transcription (when contacted with the TSPY also useful for treating prostate also useful for treating prostate cancer by inhibiting treating prostate cancer by cleaving the TSPY mRNA and therefore inhibiting its translation. The vaccine is useful for the inhibiting the development of prostate cancer in a patient
                                                                                                                                                                                                                                                                                                                                                    Diagnosing prostate cancer by determining the level of testis-specific protein Y-encoded (TSPY) mRNA or protein and comparing these TSPY mRNA or protein levels to those of a normal tissue sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
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3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nested primer 2 cloning SSH-generated 36P1A6 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 20; 32pp; English.
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99WO-US022575.
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                                                                                                                          (UROG-) UROGENESYS INC. (AFAR/) AFAR D E.
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RESULT 556

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UROG-) UROGENESYS INC.
                    WO200020584-A2.
                                                                                       02-0CT-1998;
29-JUL-1999;
Homo sapiens.
                                                                  02-OCT-1999;
                                            13-APR-2000
                                                                                                                                   (AFAR/) (HUBE/) H
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98US-0102744P.

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The human 36PlA6 gene encodes a putative transcription factor based on homology to the murine EHF gene which encodes a transcription factor which is a member of the ETS family. 36PlA6 is expressed in androgen-dependent LAPC prostate cancer xenografts and in normal prostate at approximately equal levels. The highest expression is the the prostate and colon. 36PlA6 may be involved in activating tumor-promoting genes or repressing genes that block tumorigenesis. The 36PlA6 cancer, e.g. prostate, bladder, cervical, ovarian, pancreatic and colonic cancer (all claimed). Anti-36PlA6 antibodies may be used for purifying 36PlA6 and for isolating 36PlA6 antibodies may be used for purifying 36PlA6 and for isolating 36PlA6 antibodies may be used for oligonucleotides and ribozymes can be used to inhibit the transcription and translation of the 36PlA6 gene (Claimed). The 36PlA6 polymucleotides and immunogenic fragments may also be used in cancer and polypeptides and immunogenic fragments may also be used in cancer
                                                                                                                                                                                                                                                                                                                                                                                                           Novel putative transcription factor gene 36P1A6 for treatment, diagnosis and prevention of prostate, bladder, cervical, ovarian, pancreatic, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                  Wer DE, Hubert RS, Mitchell SC;
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                                                                                                                                                                                                                                                   AFAR D E.
HUBERT R S.
MITCHELL S C.
                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-303772/26
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Gaps ; 0 Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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AAC64567 standard; DNA; 20 BP RESULT 558 AAC64567 

AAC64567;

(first entry) 14-FEB-2001 Human, prostate specific gene, 30P3C8; prostate cancer, diagnosis; cytostatic; gene therapy, vaccine; tumour; primer; ss.

Human prostate specific 30P3C8 nested primer 2 SEQ ID NO:25.

Homo sapiens

WO200061610-A2

19-OCT-2000

12-APR-2000; 2000WO-US010218 

12-APR-1999; 99US-0128860P

(UROG-) UROGENESYS INC

Saffran DC; Raitano AB, Leong K, Afar DE,

30P3C8 polypeptide and polynucleotide used for diagnosing, treating and monitoring development of prostate cancer. WPI; 2000-619224/59

Example 1; Page 57; 99pp; English.

The present invention describes human prostate specific protein 30P3CB, which is over-expressed in prostate cancer cells. 30P3CB has cytostatic cativity and can be used in vaccines and gene therapy. Methods for activity and can be used in vaccines and gene therapy. Methods for cancering the levels of 30P3CB protein or mRNA in prostate tissue, bone tissue, lymphatic tissue, serum, blood or seemen are used for diagnoshing the presence of cancer in an individual or disregulated cell growth e.g. byperplasta. The cancers which are detected or diagnosed are of the bladder, pancreas, colon, brain, bone, lung, kidney or prostate by using test samples of serum, blood or urine or tissues of the bladder, compared or sequences can be used for treating cancers expressing colymucleotide sequences can be used for treating cancers expressing colymucleotide sequences can be used for treating cancers expressing colymucleotide sequences can be used for treating cancers app3CB are used in vaccines to inhibit the development of cancer. Anti-30P3CB concorded in the development of cancer. Anti-30P3CB concorded and other proteins e.g. receptors for which 30P3CB is a ligand.

30P3CB may be a growth factor or other molecule involved in tumour growth cancer prometing activities of 30P3CB. The assays are invasion or other cancer promocing activities of 30P3CB. The assays are used for detecting, staging and monitoring prostate cancer. The 30P3CB concert and provide a more specific assay than the serum prostate cancer and provide a more specific assay than the serum prostate concert and provide a more specific assay than the serum prostate concert and provide a more specific assay than the serum prostate concert and provide a more specific assay than the serum prostate concert which is used in the exemplification of the present invention.

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps .. Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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RESULT 559 AAC64486

AAC64486 standard; DNA; 20 BP.

AAC64486;

13-FEB-2001 (first entry)

Prostate tumour associated gene 24P4C12 nested primer 2 SEQ ID NO:41. 

Human; prostate tumour associated gene; 24P4C12; prostate cancer; transmembrane protein; diagnosis; anticancer; cytostatic; vaccine; gene therapy; PCR primer; ss.

Homo sapiens.

WO200061746-A1.

19-OCT-2000.

12-APR-2000; 2000WO-US010039

The present invention provides the protein and coding sequences of human cancer related protein 20P2H8. The gene, which is found at chromosome 15q32-23, is upregulated in cancers such as that of the prostate, bladder, colon and pancreas. The sequences can be used to diagnose and treat these cancers, and to vaccinate against them. The present sequence is a PCR primer for the coding sequence of the invention

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                                                                                                                              Novel 24P4C12 polypeptides and polynucleotides, used in the diagnosis and treatment of cancer, especially prostate cancer.
                                                                                                                                                                                                           The present invention describes a prostate tumour associated gene, designated 24P4C12, and its encoded protein. 24P4C12 has anticancer and cytostatic activity, and can be used in vaccine production and in gene therapy. A pharmaceutical composition or vaccine comprising 24P4C12 can be used to treat a patient with cancer, especially prostate cancer, the vaccine can also be used to inhibit the development or progression of cancer. The polypeptides and polynucleotides can be used to diagnose cancers, especially prostate cancer. A transgenic animal comprising 24P4C12 can be used for the development and screening of therapeutic reagents. The polypeptide is a transmembrane protein which is expressed specifically in prostate cancer, allowing the development of more specific anticancer therapies and diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, cancer related protein 20P2H8; vaccine; chromosome 15g32-23; prostate cancer; bladder cancer; colon cancer; pancreatic cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hubert RS, Mitchell SC, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
                                                                Leong K, Raitano AB, Saffran DC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cancer related protein 20P2H8 cDNA PCR primer #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                Example 1; Page 65; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 gcenegreeceeceagea 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF85709 standard; DNA; 20 BP
 12-APR-1999; 99US-0128858P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 78.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Afar DEH, Raitano AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UROG-) UROGENESYS INC.
                              (UROG-) UROGENESYS INC.
                                                                Hubert RS,
                                                                                                  WPI; 2000-672681/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF85709;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                Afar DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 560
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20P2H8 polynucleotides and polypeptides useful for diagnosing and treating cancer, and for screening for screening for modulating

WPI; 2001-308645/32.

Example 1; Page 64; 111pp; English.

treating c compounds.

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The present invention relates to methods and compositions for the diagnosis and therapy of prostate cancer which utilise human SGP28 (specific granule protein 28) gene and proteins. The method involves centering cancers, particularly of prostate and colon, from cancers, particularly of prostate and colon, from coverexpression of SGP28 protein. The expression of SGP28, which is an coveracellular protein is restricted to the prostate and ovary, and is markedly up-regulated in prostate tumours. SGP28 sequence is used for diagnosis (including in vivo inaging), staging, monitoring and prognosis of prostatic and colon cancer, and for assisting salection of therapy. Composition or vaccine that contains a vector expressing an antibody specific for SGP28 protein, nucleic acid encoding SGP28 specific antibody specific for SGP28 protein, nucleic acid encoding SGP28 specific antibody optionally conjugated to toxin or therapeutic agent. SGP28 gene product is also used as source of therapeutic antisense or ribosyme agents, as primers/probes for diagnosis or prognosis, to identify compounds that inhibit calcium entry into prostatic cells, for recombinant production of SGP28 peptides and for isolating related sequences. SGP28 protein and its
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, specific granule protein 28; SGP28; therapy; PCR primer; prostate; colon; cancer; prognosis; vaccine; anticancer; SSH; suppression subtractive hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting cancers, particularly of prostate and colon, from overexpression of SGP28 protein, also methods for treating these cancers e.g. by vaccination with the protein.
                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                       3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
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                                                                                                                                  Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SGP28 gene fragment amplifying NP2 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 59; 102pp; English.
                                                                                                                                                                                                                                                   319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                       2 gcgrcgrcgcgccgaggA 20
                                                                                                                                                                                                                                                                                                                                                                                       AAD06232 standard; DNA; 20 BP.
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                                                                                                                                                                       Query Match
Best Local Similarity 78.9
Matches 15; Conservative
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Wed Apr 21 12:58:21 2004

fragments are used to raise specific antibodies (Ab) and to identify specific binding agents (potentially useful as therapeutic and diagnostic agents) and also potential anticancer agents. The present sequence is a nested primer 2 (NP2) used to amplify gene fragments resulting from SSH (suppression subtractive hybridisation) reaction. This sequence is used in the SSH isolation of cDNA fragment of human SGP28 gene 8888888888

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

0; Gaps 3.0%; Score 12.6; DB 1; Length 20; llarity 78.9%; Pred. No. 4.9e+02; Conservative 0; Mismatches 4; Indels 319 GCGTGCTGGCGGCGACGA 337 Local Similarity nes 15; Conserva Query Match Best Loc Matches

2 dedragrededeceadea 20

AAD04811 standard; DNA; 20 BP. RESULT 562 AAD04811 

AAD04811;

(first entry) 17-JUL-2001

Human 36P6D5 gene fragment amplifying primer NP2.

Human, 16P6D5 protein, secreted tumour antigen, therapy, cancer, kidney, bladder, ovary, breast, pancreas, colon, lung, vaccine, cytostatic, SSH, suppression subtractive hybridisation, PCR primer, ss.

Homo sapiens.

WO200131015-A2.

03-MAY-2001

30-OCT-2000; 2000WO-US029894.

99US-0162417P. 28-OCT-1999;

(UROG-) UROGENESYS INC

Jakobovits A, Faris M, Afar DEH, Hubert RS; Raitano AB, Mitchell SC;

WPI; 2001-308646/32.

Detecting presence of cancer expressing 36P6D5 protein in individual by comparing protein level in test sample to normal sample, where elevated level of protein in test sample indicates presence of cancer.

Example 1; Page 70; 113pp; English

The present invention relates to a gene and its encoded secreted tumour antigen, termed 36F0D5. These sequences are used for the diagnosis and treatment of various cancers which express 36F0D5, such as cancers of the kidney, bladder, ovary, breast, pancreas, colon and lungs. In normal individuals 36F0D5 protein, is predominantly expressed in pancreas, with comparising immunogenic protein of 36F0D5 is useful for inhibiting the development of prostate or colon cancer. Pharmaceutical composition comprising 3FF0D5 protein is useful for diagnosis and/or programs of prostate cancer and other cancers, for modulating or inhibiting the expression of 3FF0D5 genes and/or translation of the 3FF0D5 transcripts, and as therapeutic agents. The present sequence is a nested primer (NP)2 used to amplify gene fragments resulting from SSH (suppression subtractive hybridisation) reaction. This sequence is used in the SSH isolation of cDNA fragment of human 3FF0D5 gene

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other

Gaps ö Score 12.6; DB 1; Length 20; Pred. No. 4.9e+02; 0; Mismatches 4; Indels 3.0%; Query Match Best Local Similarity 78.9 Matches 15; Conservative

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ઠ 셤 AAF76012 standard; DNA; 20 BP

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AAF76012;

PCR primer NP2, SEQ ID NO:18, used in human PC-LECTIN cDNA isolation 22-MAY-2001

Human, PC-LECTIN; C-type lectin; transmembrane antigen; normal testis; layilin homologue; prostate cancer antigen; overexpression; androgen-dependent prostate cancer; diagnosis; prognosis; PCR primer; ss.

Synthetic.

WO200112811-A1.

22-FEB-2001

11-AUG-2000; 2000WO-US022065.

99US-0148935P. 12-AUG-1999;

(UROG-) UROGENESYS INC.

Jakobovits A, Raitano AB; Afar DEH, Hubert RS,

WPI; 2001-211222/21.

New PC-LECTIN polynucleotide encoding a transmembrane antigen over expressed in human prostate cancer, useful for the prognosis, diagnosis and treatment of prostate cancer.

Example 1; Page 59; 116pp; English

The invention relates to a novel human C-type lectin transmembrane cantigen, PC-LECTIN (AAB73309) and cDNA encoding it (AAF76004). The antigen, PC-LECTIN (AAB73309) and cDNA encoding it (AAF76004). The expression of the human PC-LECTIN gene is normally restricted to the capters of the human PC-LECTIN gene is normally restricted to the capters of the human PC-LECTIN and expression is therefore likely androgen-dependent prostate tumours. Gengard in the presente of androgen. Human PC-LECTIN therefore concer androgen-dependent prostate cancer. Human PC-LECTIN exhibits to particularly androgen-dependent prostate cancer. Human PC-LECTIN exhibits concerned to the normal particularly over a 265 residue overlap), but is not thought to be the human orthologue of layilin, as diverges significantly in a key functional domain proposed for the layilin concerned to protein. Human PC-LECTIN an immunogenic portion thereof, a vector protein. Human PC-LECTIN an immunogenic portion thereof, a vector concided-targetted ribozyme, or an anti-PC-LECTIN antibody may be used to prepare a composition for treating a patient with a cancer, concided-targetted ribozyme, or an anti-PC-LECTIN proteins are also useful for diagnosing the presence of cancer. PC-LECTIN antibodies and nucleotides are useful in the colon, pancreatic, testicular, expressing cancers. PC-LECTIN antibodies may also be used as drug targetting agents. The PC-LECTIN concludes and proteins may additionally be used in the isolation of human concerned sequence represents a pure primer used in the isolation of human concerned and proteins may additionally be used in the isolation of human concerned and proteins may additionally be used in the isolation of human concerned and proteins may additionally be used in the isolation of human concerned and proteins may additionally be used in the isolation of human concerned and proteins may additionally function or expression. 

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Gaps ; 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.98+02; tive 0; Mismatches 4; Indels Conservative Local Similarity les 15; Conserv Query Match Matches

RESULT 564 AAF8389(

AAF83890 standard; DNA; 20 BP. 

06-AUG-2001 (first entry)

Nested primer (NP)2 used in human PHOR-1 cDNA isolation.

G-protein-coupled receptor; prostate; cancer; PHOR-1; kidney; uterine; cervical; stomach; rectal; cytostatic; vaccine; cell function regulator; human; prostate homologue of olfactory receptor-1; PCR primer; se.

Homo sapiens.

WO200125434-A1.

12-APR-2001

05-OCT-2000; 2000WO-US027543.

05-OCT-1999;

(UROG-) UROGENESYS INC.

Jakobovits A, Faris M, Hubert RS; Afar DEH, Ja Raitano AB, Mitchell SC,

WPI; 2001-367230/38.

Novel gene designated PHOR-1, a G-protein-coupled receptor up-regulated in prostate cancer, useful as diagnostic marker and therapeutic target for cancers of prostate, kidney, uterus.

Example 1; Page 59; 139pp; English.

The invention relates to a novel G-protein-coupled receptor up-regulated in prostate cancer, termed PHOR-1. The encoding cDNA is contained in plantal designated plo10F2All deposited with ATCC as Accession No.PTA-112. PHOR-1 polypepides and polymuclectides are useful for diagnosing the presence of cancer, especially prostate, kidney, uterine, cervical, cancer parece by determining and comparing the level of the protein or mRNA expression in test and comparing the level of the protein or mRNA expression in test unsmall issue samples. Pharmaceutical compositions comprising PHOR-1 is useful for treating cancer. PHOR-1 proteins are useful for identifying ligands and other of generating antibodies which are useful in diagnostic, prognostic and for generating methodologies and for the treatment of prostate cancer. Cell insign methodologies and for the treatment of prostate cancer. Cell interactions mediated by PHOR-1 (The present sequence represents a primer useful in isolation of the PHOR-1 (prostate homologue of olfactory receptor

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ô Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels

319 GCGTGCTGGCGGCGGACGA 337

2 GCGTGGTCGCGGCCGAGGA 20

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RESULT 565 AAH99163

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AAH99163 standard; DNA; 20 BP.

AAH99163;

(first entry) 04-DEC-2001 Human prostate-related gene 83P5G4 cDNA nested primer #2.

83P5G4; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; ss; tumour; kidney; brain; bone; ovary; breast; pancreas; uterus; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; liver; single chain monoclonal antibody; serum; blood; urine; bladder; cervix; rectum; stomach; human; chromosome 1q31-q32. 

Homo sapiens.

WO200159115-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004426.

09-FEB-2000; 2000US-0181261P.

(UROG-) UROGENESYS INC.

Afar DEH, Challita-Eid PM, Faris M, Levin E; Jakobovits A; Hubert RS, A Mitchell SC,

WPI; 2001-514669/56.

An isolated 83P5G4-related protein useful as a diagnostic and/or therapeutic agent in multiple cancers such as prostate, bladder and bone cancer

Example 1; Page 55; 112pp; English.

The nucleic acid sequences represent the 83P5G4 gene and the primers and adaptors used to amplify 83P5G4 DNA. 83P5G4 exhibits prostate specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, testis, bladder, kidney, brain, bone, cervix, uterus, ovary, breast, pancreas, stomach, rectum, liver, colon and lung. The 83P5G4 polynucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an polynucleotide having the 83P5G4 capable of cleaving, a polynucleotide having the 83P5G4 capable of cleaving, a polynucleotide having the 83P5G4 cading sequence, are both; useful in the preparation of a composition for treating a patient with a cancer that expresses 83P5G4. The sequences can be used in diagnostic methods to monitor the level of 83P5G4 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

0; Gaps Match 3.0%; Score 12.6; DB 1; Length 20; Local Similarity 78.9%; Pred. No. 4.98+02; Local 15; Conservative 0; Mismatches 4; Indels Query Match Best Loca Matches

RESULT 566 AAS42202 ID AAS4220 XX

AAS42202 standard; DNA; 20

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prostate; colon; bladder; lung; ovarian; pancreatic; PCR primer; ss
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Best Local Similarity 78.9
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Jakobovits A;
                                                                                                                                                                                                                                                                    (UROG-) UROGENESYS INC
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                                                                                      WO200140276-A2
                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequences AAS42193-AAS42208 represent the 103P2D6 gene and the primers and adaptors used to amplify 103P2D6 DNA. 103P2D6 is not expressed in normal adult tissue but is aberrantly expressed in some forelal tissues contain and language to the prostate, testis, bladder, bone, cervix, ovary, breast, pancreas, colon and lung. The 103P2D6 DOPE, comprising a polymucleotide protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P2D6-related protein, antibody, that immunospecifically binds to an 103P2D6-related protein, and a ribozyme capable of cleaving a polymucleotide having the 103P2D6 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P2D6. The sequences can be used in diagnostic methods to monitor the level of 103P2D6 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, cytostatic, antiproliferative, vaccine, gene therapy,
six transmembrane epithelial antigen of the prostate-1; STEAP-1; cancer;
                                                                                                                            103P2D6; PCR primer; DNA adaptor; prostate; testis; foetal tissue; 8s; tumour; cancer; bone; ovary; breast; pancreas; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; serum; blood; urine; bladder; single chain monoclonal antibody; cervix; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NP2 primer used in isolation of STEAP cDNA fragment generated from SSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polynucleotide for treating and diagnosing prostate cancer is the 103P2D6 gene which encodes for 103P2D6-related proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Raitano AB, Afar DEH, Rastegar GS, Mitchell SC, Hubert RS;
Challita-Eid PM, Faris M, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
                                                                                         Juman prostate-related gene 103P2D6 cDNA nested primer #2.
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These 15; Conservative
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                                              17-DEC-2001
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                                                                                                                                                                                                                                                                                                                                 0-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD07091;
  AAS42202;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 567
AAD07091
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the present sequence is nested primer (NP2) which is used to isolate the human six transmembrane epithelial antigen of the prostate (STEAP) cDNA fragment generated from suppression subtractive hybridisation (SSH).

STEAP gene is used in gene therapy. Inhibiting the development or progression of a cancer (eg. prostate, colon, bladder, lung, ovarian and pancreatic) expressing STEAP or inhibiting the development or progression of a cancer (eg. prostate, colon, bladder, lung, ovarian and pancreatic) expressing STEAP or inhibiting growth or killing cells expressing STEAP, or inhibiting growth or killing cells expressing that comprises the vector encoding single chain monoclonal antibody that comprises the variable domains of the heavy and light chains of the monoclonal antibody that specifically binds to STEAP, such that the vector delivers the single chain monoclonal antibody coding sequence to the cancer cells and the encoded single chain monoclonal antibody is expressed intracellularly
                                                                                                                                                                                                                                                                                                                                                     New STEAP (six transmembrane epithelial antigen of the prostate) proteins, expressed in human cancers, useful for detecting and treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           84P2A9; PCR primer; DNA adaptor; prostate; testis; tissue; cancer; 88; leukaemia; tunour; kidney; brain; bone; skin; ovary; breast; pancreas; colon; lung; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prostate and testis-related gene 84P2A9 cDNA nested primer #2
                                                                                                                                                                                          Raitano AB, Saffran DC, Mitchell SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 70; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 dégrégregégégégégés 20
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06-DEC-2000; 2000WO-US033040.
                                                              99US-00455486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS11672 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-AUG-2001
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The nucleic acid sequences represent the 84P2A9 gene and the primers and adaptors used to amplify 84P2A9 DNA. 84P2A9 exhibits prostate and testis specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including leukaemia and tumours of the prostate, testis, kidney, brain, bone, skin, ovary, breast, pancreas, colon and lung. The 84P2A9 polymuclectide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 84P2A9-related protein, and a ribozyme capable of cleaving a polymucleotide having the 84P2A9 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 84P2A9. The sequences can be used in diagnostic methods to monitor the level of 84P2A9 gene products in serum, blood, urine and tissue and to thereby detect the presence of cancerous cells
                                                                                                                                                                                                                                                                          New 84P2A9 gene and its encoded protein, useful for diagnosing and treating cancer, e.g. leukemia and cancer of the prostate, testis, kidney, brain or bone, or for eliciting an immune response.
                                                                                                                                                         Levin E, Mitchell SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                         Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                   3xample 1; Page 71; 149pp; English.
                      26-JAN-2001; 2001WO-US002651.
                                                                  26-JAN-2000; 2000US-0178560P.
                                                                                                                                                           Jakobovits A, Afar DEH,
                                                                                                               (UROG-) UROGENESYS INC
                                                                                                                                                                                                                                WPI; 2001-502631/55.
                                                                                                                                                                                 Hubert RS;
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% Natch 3.0%; Score 12.6; DB 1; Length 20; Local Similarity 78.9%; Pred. No. 4.9e+02; Noservative 0; Mismatches 4; Indels 0; Gaps 319 GCGTGCTGGCGGCGACGA 337 Query Match ò

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2 ecercercececeaeaa 20

ABL50419 standard; DNA; 20 BP. (first entry) 17-JUN-2002 ABL50419; RESULT 569 ABL50419 A CAN THE STAN THE ST

Human 158P1F4 gene nested primer (NP)2 SEQ ID NO:736.

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

sapiens Synthetic, Ношо

WO200216598-A2.

28-FEB-2002.

22-AUG-2001; 2001WO-US026411.

22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P.

(AGEN-) AGENSYS INC

Levin E; Raitano AB, Afar DEH, Challita-Eid PM, Hubert RS, F Faris M, Ge W, Jakobovits A;

WPI; 2002-269357/31.

monitoring 158P1H4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158P1H4 gene products in biological sample.

Example 45; Page 116; 209pp; English.

The present invention describes a method for monitoring 158P1H4 gene

Comparison a biological sample from a patient who has or is suspected of

Exing cancer. The method comparises determining the status of 158P1H4

Gene products in a tissue sample from an individual, comparing the status

Co the status of 158P1H4 gene products in a normal sample, and

identifying the presence of aberrant 158P1H4 gene products in the sample.

Co identifying the presence of aberrant 158P1H4 gene products in the sample.

Co production. 158P1H4 polynucleotides may be used in monitoring genetic

abnormalities. The 158P1H4 proteins may be used in most cering

Co issep1H4 gene products in normal versus cancerous tissues and so

Co issep1H4 contains thenotype, in generating and characterising

Co issep1H4 polynucleotides may be used in assessing the status

Co issep1H4 proteins may be used in acsenting cancer

Co issep1H4 proteins may be used in acsenting cancer

Co issep1H4 proteins may be used in agencial and characterising

Co lissp1H4 or its particular domain, and for generating cancer

Co that bind to 158P1H4 are useful in diagnostic and

Co vaccines. Antibodies against 158P1H4 are useful in diagnostic and

Co cytocoxic T lymphocyte (CTL) or helper T lymphocyte (HTL) responses, and

contibodies are particularly useful in bladder cancer diagnostic and

contibodies are particularly useful in bladder cancer diagnostic and

contibodies are particularly useful in bladder cancer diagnostic and

Co prognostic assays, and inaging methodologies. The 158P1H4 gene also described in

Co chart bind to 158P1H4 are useful in bladder cancer diagnostic and

Concated to chromosome 8422-q23, and the 158P1H4 gene also described in

Concated to chromosome B422-q23, and the 158P1H4 gene also described in

Concated to chromosome sequence in equences used in the exemplification of the present invention

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

4; Indels 0; Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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Gaps

319 GCGTGCTGGCGGCGGACGA 337 2 dcdrdgrcdcddcddda 20 g ò

ABL50407 standard; DNA; 20 BP. 17-JUN-2002 (first entry) ABL50407; RESULT 570 ABL50407 

Human 158P1H4 gene nested primer (NP)2 SEQ ID NO:724.

Human; 158P1H4; chromosome 8q220q23, 158P1F4; chromosome 8q23; cancer; bladder cancer; immune response; cytotoxic T lymphocyte; CTL; HLA; human leukocyte antigen; helper T lymphocyte; HTL; PCR primer; adapter;

Homo sapiens.

Synthetic.

WO200216598-A2.

28-FEB-2002

22-AUG-2001; 2001WO-US026411.

22-AUG-2000; 2000US-0227098P.

rng.res

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Monitoring 158PIH4 gene products in biological sample from patient who has or is suspected of having cancer, useful for treating cancer, comprises identifying presence of aberrant 158PIH4 gene products in biological sample.
                                                                                                        Example 1; Page 69; 209pp; English
                             Challita-Eid PM, Hubert RS, 1
Faris M, Ge W, Jakobovits A;
10-APR-2001; 2001US-0282739P.
               (AGEN-) AGENSYS INC.
                                                    4PI; 2002-269357/31.
```

The present invention describes a method for monitoring 158P1H4 gene products in a biological sample from a patient who has or is suspected of products in a biological sample from a patient who has or is suspected of baving cancer. The method comprises determining the status of 158P1H4 gene products in a tissue sample from an individual, comparing the status of comparing the presence of aberrant 158P1H4 gene products in the sample. The status of 158P1H4 gene products in a normal sample, and comparing the presence of aberrant 158P1H4 sequences have cytostatic activity and can be used in vaccine production. 158P1H4 polymucleotides may be used in monitoring genetic abnormalities. The 158P1H4 proteins may be used in monitoring genetic comparing the malignant phenotype, in generating and characterising containing that bind to 158P1H4 or its particular domain, and for generating cancer vaccines. Antibodies against 158P1H4 are useful in diagnostic and products of the paticular domain, and for generating cytotoxic Tlymphocyte (CTL) or helper Tlymphocyte (HTL) responses, and simmunological respents for detecting 158P1H4 expressing cells. The antibodies are particularly useful in bladder cancer diagnostic and prognostic assays, and imaging methodologies. The 158P1H4 gene has been contended to chromosome 8q23, and the 158P1H4 gene has been contended to present invention has been located to chromosome 8q23, ABL60400 to ABL50429 and ABB95188 represent sequences used in the contendence of the present invention of the present invention

Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; 319 gcgrgcrgccgcggacga 337

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dedriedrededeceaaga 20 ABA98342 standard; DNA; 20 ABA98342; RESULT 571 ABA98342 

29-NOV-2002 (first entry) Nested primer (NP) 2.

55P4H4; cancer; immune response; ds; PCR primer.

Unidentified.

20-DEC-2001.

13-JUN-2001; 2001WO-US019246.

13-JUN-2000; 2000US-0211454P

(UROG-) UROGENESYS INC.

Levin E, Mitchell SC, Raitano AB; Hubert RS, Afar DEH, Faris M, Hube Jakobovits A;

WPI; 2002-098053/13.

Levin E;

Raitano AB, Afar DEH,

Novel isolated 55P4H4-related protein encoded by a gene over-expressed multiple cancers, useful as a diagnostic and/or therapeutic agent for cancer, preferably prostate cancer.

Example 1; Page 54; 160pp; English

This invention relates to an isolated 55P4H4-related protein encoded by a gene that is over-expressed in multiple cancers. The polypeptide is useful for inducing an immune response to an 55P4H protein, providing the protein comprises of at least one T cell or B cell epitope. The immune system cell is a B cell which generates antibodies that comprises of at least on T cell or B cell epitope. The specifically bind to the protein or is a T cell, preferably a cytotoxic T cell (TCI) which kills an autologous cell that expresses the 55P4H4 correction or a helper T cell (HTL) which secretes cytokines that correctioned which is considered useful for monitoring the presence of semilarationed which is considered useful for monitoring the presence of cancer in an individual, where the presence of elevated 55P4H4 mRNA or protein expression in the test sample relative to the normal tissue corrects in a prostate, kidney, testis, lung cervix, bone, bladder, brain or ovary tissue. The protein is useful in diagnostic assays that examine conductions associated with disregulated cell growth such as cancer and is also useful in forensic analysis of tissues of unknown origin, to treat a pathological condition characterized by the overexpression of 55P4H4 gene products in normal versus cancer use conduction assess the presence of perturbations in specific regions of the 55P4H4 gene. This sequence represents nested primer (NP) 2 used curing the method highlighted in the examples

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

.. 0 Query Match
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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0; Gaps

ABA03609 standard; DNA; 20 BP. 08-FEB-2002 (first entry) ABA03609; RESULT 572 ABA03609

Nested primer 2 used for human 34P3D7 cDNA isolation.

Human, 34P3D7; cytostatic; vaccine; gene therapy; cancer; human leukocyte antigen; HLA; major histocompatibility complex; MHC; HLA A1; HLA A11; HLA A02; HLA A24; HLA A3; HLA B35; HLA B7; primer; 88. CXXXEXEXEXXXXXXXXXXXXXXXXXXXXXX

Homo sapiens

WO200159110-A2.

16-AUG-2001.

08-FEB-2000; 2000US-0181020P. 08-FEB-2001; 2001WO-US004094

(UROG-) UROGENESYS INC.

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The invention relates to a polymucleotide, designated 34F3D7, encoding a 34F3D7-related protein, comprising a sequence of 2198 nucleotides fully defined in the specification. The presence of elevated 34F3D7 mRNA or protein expression indicates the presence of cancer occurring in prostate, bladder, Kanbey, brain, bone, cervical, uterine, ovarian, brast, pancreatic, stomach, colon, rectal leukocytes, liver, and lung pirotein, an antisense polymucleotide complementary to 34F3D7 polymucleotide, or a ribozyme capable of cleaving the 34F3D7 polymucleotide, or a ribozyme capable of cleaving the 34F3D7 polymucleotide, is useful for inhibiting the development of a cancer example demonstrating suppression subtractive hybridisation (SSH)-generated isolation of a CNNA fragment of the 34F3D7 gene
                                                                                                               New gene designated 34P3D7, encoding a tissue-specific protein highly expressed in prostate cancer, for use as diagnostic and/or therapeutic target for cancers, and for eliciting an immune response.
Challita-Eid PM, Hubert RS, Levin E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                          Example 1; Page 53; 112pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 dedredrededededa 20
Faris M, Afar DEH, Challi
Mitchell SC, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 78.99
                                                                    WPI; 2002-025689/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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0; Gaps

4; Indels

Human 125P5C8 gene PCR primer #3. AALS0002 standard; DNA; 20 BP 10-DEC-2002 (first entry) AAL50002; RESULT 573 

Human, 125P5C8, cancer, cytostatic, breast cancer, prostate cancer, bladder cancer, kidney cancer, colon cancer, ovarian cancer, PCR, primer,

Homo sapiens.

WO200272785-A2.

19-SEP-2002.

13-MAR-2002; 2002WO-US007855. 14-MAR-2001; 2001US-00809638

(AGEN-) AGENSYS INC.

Ge ⊠; Faris M, Challita-Eid PM, Hubert RS, Afar DEH, Raitano AB, Morrison RK, Morrison K, Jakobovits A;

WPI; 2002-713510/77.

New composition comprising a substance that modulates the status of 125P5C8 gene or a molecule that is modulated by 125P5C8, useful for treating or preventing cancer that expresses or over expresses 125P5C8.

Example 1; Page 68; 274pp; English

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Sequences AAS95810-AAS95820 represent the 1039388 gene and the primers and adaptors used to amplify 103938 DNA. 1039388 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 1039388 polynucleotide, its related protein and also peptide fragments of the protein are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 1039388-related protein, and a ribozyme capable of cleaving a polynucleotide having the 1039388 redains a persent with a cancer that expresses 1039388. The sequences can be used in diagnostic methods to monitor the level of 1039388 gene products in serum, blood, urine and tissue and to thereby detect the
The present invention relates to compositions comprising a substance that modulates the status of 125P5C8 or a molecule that is modulated by 125P5C8. The status of a cell that expresses 125P5C8 is modulated by composition is useful for treating cancer, particularly prostate, bladder, kidney, colon, ovary or breast cancer. The 125P5C8 protein and/or a nucleotide sequence encoding the protein is useful for immunising a mammal against cancer. The present sequence is a PCR primer shown in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          103P3E8; PCR primer; DNA adaptor; prostate; bladder; kidney; colon; lung; breast; rectum; stonach; tumour; cancer; cytoestatic; gene therapy; ss; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human; chromosome 9q13-q21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Challita-Eid PM, Raitano AB, Mitchell SC, Afar DEH;
                                                                                                                                                                                                                                                                               .
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                                                                                                                                                                                                                                       Match 3.0%; Score 12.6; DB 1; Length 20; Local Similarity 78.9%; Pred. No. 4.9e+02; es 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cancer-related gene 103P3B8 cDNA nested primer #2.
                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1, Page 55; 128pp; English.
                                                                                                                                                                                                                                                                                                                                319 GCGTGCTGGCGGCGACGA 337
                                                                                                                                                                                                                                                                                                                                                                           2 decrearedececeaeda 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS95820 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-APR-2000; 2000US-0196647P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NPI; 2002-061976/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Faris M.
                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                          Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 574
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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other; presence of cancerous cells SXS

.. 0 Length 20; 4; Indels Score 12.6; DB 1; Pred. No. 4.9e+02; 0; Mismatches 4; Query Match Best Local Similarity 78.9%; Matches 15; Conservative

à g 575 RESULT

AAS99443 standard; DNA; 20 BP. AAS99443

AAS99443;

12-MAR-2002 (first entry)

cancer related protein 98P7C3 nested PCR primer 2.

Human; 98P6C3; ss; homeodomain protein; vaccine; cytostatic. epitope; transgenic animal; immunogen; T cell; B cell; cytotoxic T cell; CTL; prostate cancer; bladder cancer; kidney cancer; lung cancer; breast cancer; uterine cancer; cervical cancer; stomach cancer; rectal cancer; stomach cancer; suppression subtractive hybridisation; SSH. 

Homo sapiens.

WO200190157-A2.

29-NOV-2001.

24-MAY-2001; 2001WO-US017495.

24-MAY-2000; 2000US-0207138P.

(UROG-) UROGENESYS INC.

New isolated 98P7C3-related homeodomain protein highly expressed in various cancers, useful in cancer vaccines and for generating immune response directed to 98P7C3 in mammal. Challita-Eid PM, Hubert RS, Mitchell SC, Jakobovits A; WPI; 2002-097642/13.

Levin E;

Faris M, Afar DEH,

Example 1; Page 53; 155pp; English.

the invention relates to an isolated 98P7C3-related protein which is a homeodomain protein highly expressed in various cancers. Also include are polymuclectides encoding the protein or proteins 90% identical to 98P7C3, polymuclectides encoding the protein or proteins of identical to 98P7C3, an ancient of the protein comprising the polymuclectides (including an expression vector comprising the yestor, an anti-98P7C3 anti-body, an non-homen transformed with the vector, an anti-98P7C3 anti-body, an non-homen transformed with the vector, an anti-98P7C3 anti-body, an non-homen transformed with the vector, an anti-98P7C3 anti-body, an non-homen transform or polymuclectides in a biological sample, monitoring the 98P7C3 protein or polymuclectides in a biological sample, monitoring the presence of cancer in an individual by detecting an elevated level of the 98P7C3 protein or polymuclectides and a pharmaceutical composition comprising a modulator of 98P7C3. 98P7C3 protein, or T cell/B cell comprising the protein. Upon contact with a cytotoxic T cell (CTL) the immunogen may be a nucleic acid ancoding the protein or epitope. The anti-body is useful for delivering a cytotoxic agent to a cell that expresses 98P7C3, by conjugating the conjugate. The modulator is useful for the anti-body-agent to a cell that expresses 98P7C3, by conjugating the conjugate. The modulator is useful for treating a patient with a cancer

that expresses 98P7C3 (e.g. prostate cancer, bladder cancer, kidney cancer, lung cancer, breast cancer, uterine cancer, cervical cancer, estomach cancer, rectal cancer and colon cancer), by administering to the patient a victor that comprises the modulator, such that the vector delivers a single chain monoclonal antibody coding sequence to the cancer cells and the encoded single chain antibody is expressed intracellularly in it. The gene for 98P7C3 is located on human chromosome 4q11-q12. The present sequence is oligonucleotide adapter or PCR primer used to isolate a CDNA sequence. hybridisation, SSH

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Gaps

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

; 0 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels 3.0 Best Local Similarity 78.9 Matches 15, Conservative

319 GCGTGCTGGCGGCGACGA 337 2 ecercercececcaaca 20 ઠે 셤

RESULT 576 ABK67422 ID ABK6

ABK67422 standard; DNA; 20 BP.

ABK67422;

(first entry) 02-JUL-2002 Human 83P2H3 cDNA isolation nested PCR primer 2.

Human; human leukocyte antigen; HLA; immunogen; 83P2H3; CaTrF2El1; calcium transport protein; cancer; prostate cancer; cytostatic; chromosome 7q34; chromosome 12q24.1; T cel1; B cel1; Bs; primer.

Homo sapiens.

WO200214361-A2.

21-FEB-2002.

17-AUG-2001; 2001WO-US025782.

17-AUG-2000; 2000US-0226329P.

(AGEN-) AGENSYS INC.

, Faris M, Saffran DC, Afar DEH; Jakobovits A; Raitano AB, Challita-Eid PM, Levin E, Hubert RS, Ge W,

WPI; 2002-269179/31.

Monitoring 83P2H3 gene products for monitoring the presence of cancer is a subject, comprises determining the status of 83P2H3 gene products in tissue sample from the subject and comparing it to a normal sample. 

Example 1; Page 76; 270pp; English.

The invention relates to monitoring 83P2H3 (a calcium transport protein whose gene is located on chromosome 7q34) gene products in a biological sample from an patient who has or is suspected of having cancer cancer; captured of laving cancer cancer; captured in a tissue sample from an comparing the status of 83P2H3 gene products expressed by calls in a tissue sample from an product and (b) comparing the status to the status of 83P2H3 gene products in a normal sample. Also included are modulators of 83P2H3 gene products in a normal sample. Also included are modulators of 83P2H3 function or status, generating antibodies/immune response against 83P2H3 (or related protein Carrezzil whose gene is located on chromosome 12q24.1) using identified HIA (human leukocyte antigen) binding peptides derived from the protein, delivering a cytotoxic agent to a cell. Captured from the protein antigen-binding region of the antibody, a recombinant protein comprishing an antigen-binding region of the antibody, a non-human transgenic animal that produces the recombinant protein, a

thybridoma that produces the recombinant protein, a single- chain monoclonal antibody that comprises the variable domains of the heavy and monoclonal antibody that comprises the variable domains of the heavy and light chains of the anti-aPSPH3 antibody, a vector comprising a polynucleotide that encodes the monoclonal antibody and inducing an immune response to a 83PZH3 protein, by providing a 83PZH3-related protein that comprises a T cell or B cell epicope, and contacting the epitope with an immune system T cell or B cell, respectively. The method is useful for monitoring 83PZH3 gene products in a biological sample for compitoring the presence of cancer in an individual. The modulator is useful for inhibiting the growth of cancer cells that express 83PZH3. for treating an immune response against 83PZH3, and for detecting the cancer that expresses 83PZH3. The immunological methods are useful for generating an immune response against 83PZH3, and for detecting the presence of 83PZH3-related protein or polynucleotide in a biological sample from a patient who has or who is suspected of having cancer. The semicologies and treatment, to detect and quantify 83PZH3 and mutant methodologies and treatment, to detect and quantify 83PZH3 and mutant in solution is supported brotein. For itsolating 83PZH3 calated protein of cDNA encoding 83PZH3 or its 

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

ö 0; Gaps Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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RESULT 577

ABK70514 standard; DNA; 20

.5-JUL-2002 (first entry) ABK70514;

Human cDNA 85P1B3 nested PCR primer 2.

Human; cytostatic; 85PlB3; cancer; immunogen; 88; primer; PCR; chromosome 15q14.

Homo sapiens.

WO200218578-A2.

07-MAR-2002

28-AUG-2001; 2001WO-US026838.

28-AUG-2000; 2000US-0228432P.

(AGEN-) AGENSYS INC.

Challita-Eid P; Ge ₩, Faris M, Hubert RS, Afar D, Raitano AB, F Jakobovits A;

WPI; 2002-382963/41.

Composition for modulating the status of 85P1B3 protein or a molecule comprising a substance e.g. antibody specific to, nucleic acid encoding, or ribozyme of 85P1B3.

Example 1; Page 76; 201pp; English.

The invention relates to a composition comprising a substance that modulate the status of 85P1B3, where the status of a cell expresses 85P1B3 gene product is modulated. Also included are a composition 

comprising a peptide region of 5 amino acids of the 85PBB protein, in selected from an apposition having a value greater than 0.5 in the selected from an apposition having a value greater than 0.5 in the professible research beaving a value less than 0.5 in the percent accessible research baving a value greater than 0.5 in the percent accessible research by the profile, or an as position having a value greater than 0.5 in the percent accessible research by the profile, or an as position having a value greater than 0.5 in the beta-turn profile; a polynucleotide that encodes analogue peptide of 8, 9, 10 or 11 contiguous cessions of a monoclonal antibody; a non-human transgenic antigen-binding region of a monoclonal antibody; a non-human transgenic antigen-binding region of a monoclonal antibody; a non-human transgenic confer comprises antibody specific to the protein; a single chain monoclonal antibody (MAD) that conforts to the protein; a single chain monoclonal antibody (MAD) that conforts the protein; a vector comprising a polymucleotide that encodes the MAD; inhibiting growth of comprising a polymucleotide the protein, antisense polymucleotide of the protein, and generating a mammalian immune response the protein, and generating the protein, and generating a mammalian immune specifically recognize the protein, and generating a mammalian immune specifically recognize the protein, and generating a mammalian immune specifically recognize the protein, and generating a mammalian immune specifically crecognize the protein, and generating a mammalian immune specifically crecognize the protein, and generating a mammalian immune specifically crecognize the protein, and generating a mammalian immune specifically crecognize the protein, and generating a mammalian immune specifically crecognize the protein, and generating a mammalian immune specifically crecognize the protein, and generating a protein, generating a cytoroxic agent conjugated to a meniod are useful for inhibiting concer cells or treating a mammalian immun

Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps .. Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels

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AAL40496 standard; DNA; 20 BP.

AAL40496;

19-SEP-2002 (first entry)

158P1D7 cDNA related PCR primer SEQ ID No 668.

Cytostatic; 158P1D7; cancer; bladder cancer; mouse; rat; rabbit; dog; cat; cow; horse; human; vaccine; gene therapy; PCR; primer; ss. 

Homo sapiens.

WO200216593-A2.

28-FEB-2002.

22-AUG-2001; 2001WO-US026276.

22-AUG-2000; 2000US-0227098P. 10-APR-2001; 2001US-0282739P.

(AGEN-) AGENSYS INC

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Levin
Raitano AB, Afar DEH,
Faris M, Hubert RS, Raitano A
Challita-Eid PM, Jakobovits A;
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WPI; 2002-425659/45.

ö New compositions comprising a gene (designated 158P1D7), its encoded protein or their modulators, useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. dogs, cats, cows, horses humans)

Example 1; Page 68; 181pp; English

The invention relates to a novel nucleic acid, designated 158PID7. The compositions are useful for treating or diagnosing cancers, particularly bladder cancer, in mammals (e.g. mice, rats, rabbits, dogs, cats, cows, horses or humans). The compositions are also useful for monitoring genetic abnormalities and in preparing cancer vaccines. The nucleic acid of the invention can be used in gene therapy to treat the said disorders. This polynucleotide sequence represents a PCR primer of the 158PID7 cDNA invention

Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels 15; Conservative Local Similarity Query Match Matches

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RESULT 579

AAL53476 standard; DNA; 20 BP AAL53476

(first entry) 16-JAN-2003 AAL53476;

Zinc transporter protein 108P5H8 nested primer 2.

Cytostatic; gene therapy; vaccine; zinc transporter protein 108P5H8; cancer; breast; colon; ovarian; lung; humoral; cellular immune response; passive immunisation; PCR; primer; ss. 

Unidentified

WO200260953-A2

17-DEC-2001; 2001WO-US049133.

15-DEC-2000; 2000US-0256210P

(AGEN-) AGENSYS INC.

Mitchell SC; Hubert RS, N Jakobovits A; Eid PM, Faris M, Afar DEH, Morrison KJM, Raitano AB, Challita-Eid PM, Levin E,

WPI; 2002-627469/67.

Composition comprising a substance that modulates the status of a zinc transporter protein (108P5H8), useful in diagnosing and treating patients with cancer that express 108P5H8, such as breast, colon, ovarian or lung

Example 1; Page 95; 309pp; English.

The invention relates to a new composition comprising a substance that modulates the status of a zinc transporter protein, designated as 108P5H8, or a molecule that is modulated by 108P5H8. The composition is

useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 108P5H8, such as breast, colon, ovarian or lung cancer. The 108P5H8 gene or its fragment can be used to elicit humoral or cellular immune response. The antibodises are useful in active or passive immunisation. The 108P5H8 polymucleotides are useful as probes and primers for the amplification or detection of 108P5H8 genes, as coding sequences for directing the expression of 108P5H8 polypeptides, or as tools for modulating or inhibiting the expression of 108P5H8 genes. The polymucleotides of the invention can be used to treat disorders by gene therapy. This polymucleotide sequence represents a zinc transporter protein 108P5H8 related PCR primer of the invention

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Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;

Gaps ö 3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ive 0; Mismatches 4; Indels Query Match
Best Local Similarity 78.9
Matches 15; Conservative

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RESULT 580 ABV99876

ABV99876 standard; DNA; 20 BP.

ABV99876;

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(first entry) 28-MAR-2003 Human 121P2A3 post-SSH nested PCR primer 2.

Human, 121P2A3; cytostatic; immunostimulant; vaccine; PCR; primer; humoral immune response; cellular immune response; 88; suppression subtractive hybridisation; SSH.

Ношо

WO200283068-A2.

24-OCT-2002

10-APR-2001; 2001US-0282739P. 25-APR-2001; 2001US-0286630P. 22-JUN-2001; 2001US-0300373P. 09-APR-2002; 2002WO-US011359.

(AGEN-) AGENSYS INC

Mitchell SC; W, Jakobovits A; Challita-Eid PM, Raitano AB, Faris M, Hubert RS, Afar DEH, Saffran D, Morrison K, Morrison RK, Ge WPI; 2003-092956/08. 

New composition comprising a substance that modulates the status of 121P2A3 polypeptides, useful for eliciting humoral or cellular immune responses or in assessing the status of 121P2A3 gene products in normal versus cancerous tissues.

Example 1; Page 70; 362pp; English.

The invention relates to a novel composition comprising a substance that modulates the status of a protein, 121P2A3. The composition of the invention has cytostatic and immunostimulant activity, and is useful as a vaccine. The 121P2A3 proteins and polynucleotides are useful for eliciting humoral or cellular immune response. The polynucleotides are useful for characterising cytogenetic abnormalities of this chromosomal locus, as tools that can be used to delineate cytogenetic abnormalities in the chromosomal region that encodes 121P2A3 that may contribute to malignant phenotype, and in assessing the status of 121P2A3 gene products in normal versus cancerous tissues. The proteins are useful for

ABT43860 standard; DNA; 20 BP

RESULT 582

(first entry)

16-OCT-2003

ABT43860;

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generating and characterising domain-specific antibodies, for identifying agents or cellular factors that bind to 121P2A3 or a particular structure domain, and in various therapeutic and diagnostic contexts, including cancer vaccines. The antibodies or T cells reactive with the product are useful in passive or active immunisation, and in imaging methodologies bor the antanagement of cancer. The present sequence represents an nested PCR primer used in the invention to amplify gene fragments resulting from suppression subtractive hybridisation (SSH) reactions
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Ge W, Raitano AB, Challita-Eid PM;
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25-APR-2001; 2001US-0286630P.
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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The invention relates to a novel composition comprising a substance that modulates the status of a 1519304 protein (e.g. 1519304 variant 1-11; or a molecule that is modulated by the 1519304 protein; where the status of a cell that expresses the 1519304 protein is modulated. The novel compositions, or the 1519304 protein and genes, are useful for eliciting a humoural or cellular immune response. The 1519304 genes and proteins are also useful for diagnosing, proventing preventing or treating cancer, e.g. adenocarcinoma, bladder cancer, colorectal cancer, lung or bronchial cancer, breast cancer or carcinoma. This polynucleotide sequence represents a 1519304 related primer of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New 151P3D4 proteins and genes, useful for eliciting a humoral or cellular immune response, or for diagnosing, prognosing, preventing or treating cancer, e.g. adenocarcinoma, bladder cancer, lung, breast cancer
                                                                                                             Cytostatic; gene therapy; vaccine; modulator; 151P3D4; humoural; cancer; cellular immune response; adenocarcinoma; bladder; colorectal; lung; bronchial; breast; carcinoma; PCR; primer; ss.
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Morrison RK, Ge W, Jakobovits A;
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                                                                          DPNCDN nested primer 2 (NP2).
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25-APR-2001; 2001US-0286630P.
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                                                                                                                                                                                        Unidentified.
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Homo sapiens.

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The invention relates to a novel composition comprising a substance that modulates the status of a 162P1B6 protein. The protein comprises one of 1 status of a 162P1B6 protein. The protein comprises one of 2 sequences of 70 - 146 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of a cell that expresses the protein is modulated. An antibody to the 162P1B6 protein is used to deliver a cytotoxic agent or a diagnostic agent to a cell that expresses the 162P1B6 protein. The composition is used to inhibit the growth of cancer cells or generate an immune response. The composition is used for detecting the presence of a 162P1B6 related protein or a 162P1B6 related polymucleotide in a sample. The 162P1B6 proteins and polymucleotides encoding them are useful for diagnosing, proteins or reating cancer, or breast cancer. They can also be used for eliciting a humoral or cellular immune response. The artibodies or T cells reactive with 162P1B6 are useful for active or passive immunisation. Transgenic animals are useful for developing and screening of useful reagents. The polymucleotide and polypeptide screening of the invention can also be used to treat disorders by being used in a vaccine or in gene therapy. This polymucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                           Composition for diagnosing, prognosing, preventing or treating cancer, for eliciting a humoral or cellular immune response, or for active or passive immunization, comprises a substance that modulates the status of a 162P1E6 protein.
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transgenic animal; vaccine; gene therapy; PCR; primer; ss.
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2001US-0286630P.
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25-APR-2001;
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The invention describes a composition comprising a substance that modulates the status of a protein (1) of 340 or 283 amino acids, or of any of the 15 sequences of 259 amino acids, given in the specification, or a molecule that is modulated by the protein, where the status of the cell that expresses the protein is modulated. The compositions, proteins, polymucleotides and methods are useful for generating and electring cancer. The STEAP-1-related proteins are useful for generating cancer vaccines. The polymucleotides are useful as tools for delineating, with greater precision, cytogenetic abnormalities in the chromosomal region that encodes STEAP-1 that may contribute to the malignant phenotype. This sequence represents a primer used to analyse human six transmembrane epithelial antigen of the prostate or STEAP-1 CDNA's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                             당
                                                                                                                                                                                                                                                                            STEAP-1-related protein, useful for treating and detecting cancer.
                                                                                                                                                                                                                                                             status
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; vaccine; cancer; immune response; PCR; primer; ss.
                                                                                                                                                                                               Ge W, Raitano AB, Challita-Eid PM, Jakobovits A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hubert RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
3.0%; Score 12.6; DB 1; Length 20;
Best Local Similarity 78.9%; Pred. No. 4.9e+02;
Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                           New composition comprising a substance that modulates the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Faris M,
                                                                                                                                                                                                                                                                                                           Example 1; Page 70; 248pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       319 GCGTGCTGGCGGGGGGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jakobovits A, Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 ecercercececceacea 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ78176 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-APR-2001; 2001US-0282739P.
10-APR-2001; 2001US-0283112P.
25-APR-2001; 2001US-0286630P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-APR-2002; 2002WO-US011654.
                                                                                       06-SEP-2002; 2002WO-US028371.
                                                                                                                     06-SEP-2001; 2001US-0317840P. 05-APR-2002; 2002US-0370387P.
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                                                                                                                                                                                                                               WPI; 2003-313240/30.
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                                                                                                                                                                   (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nested primer #2
                            WO2003022995-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200283921-A2
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                                                           20-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ78176;
                                                                                                                                                                                                   Faris M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 585
ABZ78176
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The present invention relates to novel human cancer-related genes and proteins (ABZ78120-ABZ78168 and ABR01789-ABR01861). The genes and proteins are useful for eliciting a humoral or cellular immune response. The genes are useful for probes and primers for the amplification and/or detection of genes, mRNAs or their fragments, as reagents for the diagnosis and/or prognosis of cancer, as coding sequences capable of directing the expression of the protein, as tools for modulating or inhibiting the expression of genes and/or translation of transcripts, and as therapeutic agents. The proteins and peptides are useful as therapeutic, prognostic and diagnostic reagents for cancer. The present sequence is a primer, used in an example from the invention
                                                              New composition comprising a substance that modulates the structure of proteins and polynucleotides, useful for therapeutic, prognostic and diagnostic reagents for eliciting cellular or humoral immune response in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cancer associated coding sequence; cancer; human; cytostatic; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                         3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cancer associated coding sequence PCR primer #3.
Morrison K, Morrison RK, Raitano AB;
                                                                                                                                                 Example 1; Page 72; 1021pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           319 GCGTGCTGCCGCCGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GCGTGGTCGCGGCCGAGGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-APR-2001; 2001US-0282739P.
10-APR-2001; 2001US-0283112P.
25-APR-2002; 2001US-0286630P.
10-APR-2002; 2002US-00286630.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ20563 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-APR-2002; 2002WO-US011645.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33-MAR-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            15; Conservative
                               WPI; 2003-075555/07.
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                  cancer patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200283920-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ20563;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 586
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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New pharmaceutical composition for diagnosing, prognosing, preventing or treating cancer, comprises a substance that modulates a nucleic acid sequence, e.g. 105P1B7, 152P1A2B or 156P3A6, or a molecule modulated by the nucleic acid.

Example 1; Page 34; 72pp; English.

Challita-Eid PM;

Jakobovits A, Hubert RS,

(AGEN-) AGENSYS INC

WPI; 2003-093030/08.

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The present invention relates to a pharmaceutical composition comprising a substance that modulates the status of a cancer associated nucleic acid sequence such as given in the specification (see ABZ20564-ABZ20575) or a molecule that is modulated by the above nucleic acid sequence, where the status of a cell that expresses the nucleic acid sequence is modulated. The composition is useful in diagnosing, prognosing, preventing and/or treating cancer. The nucleic acid sequence may be used in monitoring antibodies, for identifying agents or cellular factors that bind to a protein, and in therapeutic and diagnostic contexts, such as diagnostic assays, cancer vaccines, and methods of preparing vaccines. The present sequence is a primer used to identify the cancer associated coding sequences suitable to be modulated in the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises the amino acid and coding sequence of a 184PIE2 protein. The DNA and protein sequences of the invention are useful for diagnosing, prognosing, preventing and/or treating cancer. The 184PIE2 DNA and protein sequences may also be used to elicit a humoral or a cellular immune response in patients and in monitoring generic abnormalities. Antibodies raised against the 184PIE2 proteins may be use in active or passive immunisation. The present DNA sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New 184PIE2 polynuclectide encoding a 184PIE2 protein, useful for diagnosing, prognosing, preventing or treating cancer, in eliciting an immune response, and in chromosome mapping.
                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Faris M, Hubert RS, Morrison K;
                                                                                                                                                                                                                                                                                                                                                            4; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene therapy, vaccine, 184P1E2, cancer, genetic abnormality, cellular immune response, immunisation, PCR, primer, ss.
                                                                                                                                                                                                                                                                                                                        3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02;
                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               184P1E2 gene-specific nested PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 69; 394pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chalitta-Eid PM, Raitano AB, rarr
                                                                                                                                                                                                                                                                                                                                                                                                    319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GCGTGGTCGCGGCCGAGGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-APR-2002; 2002WO-US011643.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-APR-2001; 2001US-0282739P.
25-APR-2001; 2001US-0286630P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AALS2254 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-148269/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified.
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                                                                                                                                                                                                                                                                                                                          Query Match
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RESULT 589

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This invention relates to a novel gene designated 205P1B5, and the encoded protein, which is aberrantly expressed in prostate cancer. Specifically, it refers to the two variants of 205P1B5 mapped to chromosome 821-8912, namely 205P1B4v1 and 205P1B5v2 and fragments thereof that serve as useful diagnostic, prophylactic, prognostic and/ or therapeutic targets for protate and other types of cancers. The present invention describes methods for the isolation of 205P1B5, for generating an immune response and for generating pransgenic or knock out animals for the development and screening of therapeutically useful reagents. Purthermore, it refers to identifying proteins, small molecules or other agents that interact with 205P1B5, and can be used to identify pathways activated by 205P1B5. Accordingly, these are cytostatic and immunogenic compositions that are useful for the development of cancer vaccines. This suppressive subtractive hybridisation (SSH) to isolate the 205P1B5 CDNA fragment of the invention.
                                                                                                                                                                                                                                                                                                                                                               205P1B5; prostate cancer; immune response; transgenic; knock out animal; cytostatic; immunogenic; vaccine; ss; SSH; suppressive subtractive hybridisation; PCR; primer; NP2.
                                                                                                                                                                                                                                                                                                                              Nested PCR primer 2 (NP2) used in SSH to isolate 205P1B5 cDNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotide designated 205P1B5, for diagnosing and treating prostate cancer, and as probes or primers for the amplification and/or detection of 205P1B5 genes.
                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Raitano AB, Faris M, Hubert RS, Jakobovits A;
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3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; ative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 60; 162pp; English.
                                                                          319 GCGTGCTGGCGGCGACGA 337
                                                                                                              2 dégrégredégécekadá 20
                                                                                                                                                                                                           ADC71183 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-AUG-2002; 2002WO-US027760.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-AUG-2001; 2001US-0316664P.
                                                                                                                                                                                                                                                                                    18-DEC-2003 (first entry)
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Query Match
Best Local Similarity 78.9%
Matches 15, Conservative
              Local Similarity 78.9 tes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003020954-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-MAR-2003.
                                                                                                                                                                                                                                                   ADC71183;
    Query Match
                                                                                                                                                                      RESULT 588
                                       Matches
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The present invention describes a composition (I) comprising a substance that modulates the status of 121PIFI (gene and encoded protein), or a molecule that is modulated by 121PIFI (gene and encoded protein), or a molecule that is modulated. The human 121PIFI gene maps to chromosome 40. (I) has cytostatic activity, and can be used in gene therapy, and in vaccines. The composition (I) can be used for diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIFI, such as breast, colon, ovarian or lung cancer. The 121PIFI gene or its cragment can be used to elicit a humoral or cellular immune response. I21PIFI antibodies can be used in active or passive immunisation. 121PIFI polymolecides are useful as probes and primers for the amplification or detection of 121PIFI genes, as coding sequences for the amplification or inhibiting the expression of 121PIFI genes. The present sequence is used in the exemplification of 121PIFI genes.
                                                                                                                                                     121P1F1; 121P1F1 modulation; human; chromosome 4q; cytostatic; gene therapy; vaccine; cancer; immune response; immunisation; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Composition comprising a substance that modulates the status of 1219171, useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 1219171, such as breast, colon, ovarian or
                                                                                                                                                                                                                                                                                                                                                                                                                                                Ge ⊠;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
                                                                                                                                                                                                                                                                                                                                                                                                                                              Challita-Eid PM, Hubert RS, Raitano AB, Faris M, Afar DEH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                      121P1F1 gene nested primer (NP) 2 SEQ ID NO:721.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 71; 285pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 dedregrededededada 20
           ADD84533 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                     28-FEB-2002; 2002WO-US006242.
                                                                                                                                                                                                                                                                                                                                                                        05-MAR-2001; 2001US-00799250.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE65924 standard; DNA; 20
                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-156757/15.
                                                                                                                                                                                                                                                                                                                                                                                                           (AGEN-) AGENSYS INC.
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                              WO200295009-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jakobovits A;
                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lung cancer.
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                                                                                 29-JAN-2004
                                                                                                                                                                                                                                                                                                   28-NOV-2002.
                                                                                                                                                                                                             Synthetic.
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                                                ADD84533;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 590
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ADD8453
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319 GCGTGCTGGCGCCGGACGA 337

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15-MAY-2003

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This invention relates to novel composition comprising a substance that modulates the status of a 433 residue protein, given in the specification with the DNA sequence encoding it, or a molecule that is modulated by the protein. The novel protein 193PEIB exhibits tissue specific expression in normal adult tissue and is aberrantly expressed in cartain cancers. Compositions which modulate the 193PIBIB protein may have cytostatic activity and the DNA sequence which encodes protein 193PIBIB may be useful in gene therapy. The composition of the invention may be useful por the treatment of cancer. The present sequence is that of an RT-PCR primer which was used for the amplification of human 193PIBIB gene DNA during the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Typing nucleic acid for obtaining information about several variable sites involves simultaneously or sequentially performing two or more primer extension reactions, and determining the pattern of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single nucleotide polymorphism; nucleic acid typing; tissue typing; human; PCR; primer; angiotensin converting enzyme; ACE; ss.
                                                                                                                                                                                             New composition comprising 193PIE18-related protein, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Angiotensin converting enzyme SNP fragment Eu6 PCR primer B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.0%; Score 12.6; DB 1; Length 20; Best Local Similarity 78.9%; Pred. No. 4.9e+02; Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                  Hubert RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                    Faris M,
                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 59; 260pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pourmand N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PYRO-) PYROSEQUENCING AB.
(STRD ) UNIV LELAND STANFORD JUNIOR.
(GARD/) GARDNER R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          319 GCGTGCTGGCGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 degregregedegada 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 592
ABN79977/c
ID ABN79977 standard; DNA; 14 BP.
                                                                                                                                                                                                                       preventing or treating cancer.
                                                                                    Challita-Eid PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-2001; 2001WO-GB004042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8-SEP-2000; 2000GB-00022069
07-DEC-2001; 2001US-00013312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ronaghi M, Ekstroem B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PI; 2002-393849/42
                                                                                                                                                      WPI; 2003-532905/50.
                                         (AGEN-) AGENSYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200220837-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-MAR-2002
                                                                                    Raitano AB,
Jakobovits A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABN79977;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a novel composition which comprises a substance that modulates the status of a novel protein (1612210B) and its variants having a sequence of 875 amino acids provided in the specification. The protein of the invention is over-expressed in certain cancers. The compounds of the invention may have cytostatic activity and the sequence of the 1612210B protein, and the gene which encodes it, may be useful for gene therapy or the development of a vaccine. The composition and methods of the invention are useful in diagnosing, preventing and useful careating ancier. The present sequence is that of PCR primer which was used for amplification of a region of the gene encoding the human 16122F10B protein during the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      193P1E1B; tissue specific expression; cancer; cytostatic; gene therapy; cancer; human; PCR; RT-PCR; reverse transcription PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A composition for diagnosing, preventing and treating cancer (e.g. prostatic, renal or uterine cancer) comprises 161P2F10B polynucleotides and polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                      Jakobovits A, Raitano AB, Faris M, Hubert RS, Ge W, Morrison KJM;
Morrison RK, Challita-Eid PM;
                                           161P2F10B; cancer; cytostatic; gene therapy; vaccine; PCR; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.0%; Score 12.6; DB 1; Length 20; 78.9%; Pred. No. 4.9e+02; tive 0; Mismatches 4; Indels
Human 161P2F10B protein-related PCR primer SeqID36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Juman protein 193P1E1B-related PCR primer SeqID59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 36; 135pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            319 GCGTGCTGGCGGCGGACGA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 GCGTGGTCGCGGCCGAGGA 20
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                                                                                                                                                                                                                                                                                           07-NOV-2001; 2001US-00005480.
                                                                                                                                                                                                                                                  17-NOV-2002; 2002WO-US036002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD96944 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-441560/41.
                                                                                                                                                                                                                                                                                                                                                             (AGEN-) AGENSYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003050255-A2
                                                                                                                                                        WO2003040340-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JAN-2004
                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUN-2003
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incorporation

06-DEC-2002; 2002WO-US039274

ADD96944;

RESULT 591

Query Match Best Loca Matches

à 셤

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Gaps

0

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diagnosis; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel method for obtaining typing information about several variable sites within target nucleic acid, or typing one or mote nucleic acid molecules. The methods of the invention are useful for typing one or more nucleic acid molecules containing two or more variable sites, preferably nucleic acid molecules containing three or more variable sites are typed, where three or more primer exension reactions or performed. The method is also useful for diagnosis of pathological conditions characterized by the presence of specific nucleic acid molecule(s). The method are particularly suited for identifying microbial species or their subtypes, and in typing procedures e.g. typing of polymorphisms, tissue typing or in clinical applications. The sequence cucleotide polymorphism (SNP) Euc from the angiotensin converting enzyme (ACE) gene. The primer binds to the template with its 3, end 5 cnucleotides from the SNP position
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-ab; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; Philadelphia chromosome; inflammation; autoimmune disease; atheroselerosis; myocafdial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocafdial ischaemia; Kawasaki disease; septic shock; HIV; human immunodeficiency virus; acquired immune deficiency syndrome; AIDS;
                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human relA hammerhead ribozyme target sequence (nt. position 1058).
                                                                                                                                                                                                                                          Similarity 92.9%; Score 12.4; DB 1; Length 14; Similarity 92.9%; Pred. No. 2.5e+02; 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                   Sequence 14 BP; 1 A; 5 C; 4 G; 4 T; 0 U; 0 Other;
Disclosure; Fig 5B; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94US-00201109.
94US-00218934.
94US-00222795.
94US-00227958.
94US-00227958.
94US-0021919.
94US-00291932.
94US-0029183.
                                                                                                                                                                                                                                                                                                                                                                            AAT55127 standard; RNA; 15 BP.
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                                                                                                                                                                                                                                                                                          268 ACCTGGAGCAGGC 281
                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003 (revised)
21-APR-1997 (first entry)
                                                                                                                                                                                                                                          Query Match 2.9 Best Local Similarity 92.9 Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                 14 Accredadedade 1
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19-AUG-1994;
02-SEP-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-FEB-1995;
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04-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                     AAT55127;
                                                                                                                                                                                                                                                                                                                                                       RESULT 593
AATS5127/c
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The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves relA mRNA at the nucleicide abse position indicated in the DE line. The relA gene product is a subunit of the transcriptional regulator NF-kappaB and is implicated specifically in the induction of inflammtory responses. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes are designed to cleave the target cueful for treating rheumatoid arthritis, restenosis and asthma as well as in increasing observed to transplanted tissues. The procential immunosuppressive properties of a ribozyme that cleaves relA mRNA means that uses are limited to local delivery, acute indications or ex vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arthritic condition, graft tolerance, immune response; target; cleavage; hammerhead ribozyme; hairpin ribozyme; human; rebbit; mouse; collagenase; stromelysin; synovial membrane; joint; arthritis; osteoarthritis; rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stinchcomb DT, Chowrira B, Direnzo A, Draper KG, Dudycz LW; Grimm S, Karpelsky A, Kisich K, Matulic-Adamic J, Mcswiggen JA; Modak A, Pavco P, Balgleman L, Sullivan SM, Sweedler D, Thompson JD; Tracz D, Usman N, Wincott FE, Woolf T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ribozymes having modified bases and methods for producing them - for use in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; Page 229; 407pp; English.
94US-00303039.
94US-00311486.
94US-00311749.
94US-00316771.
94US-0031693.
94US-00334847.
94US-00334847.
94US-0033568.
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                                                                                                                                                                                                                                                                                                                                                              94US-00363233
                                                                                                                                                                                                                                                                                                                                                                                          95US-00380734
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-JUL-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-351090/45.
                                                                                                                           03-OCT-1994;
11-OCT-1994;
11-OCT-1994;
04-NOV-1994;
10-NOV-1994;
16-DEC-1994;
23-DEC-1994;
                                         23-SEP-1994;
23-SEP-1994;
28-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                          30-JAN-1995;
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neutral lipid transfer; plasma lipoprotein; atherosclerosis; athereccomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolosemia; dyslipidaemia; hypoalphalipoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angloplastic restenosis; low density lipoprotein; diabetes; HDL; human;
Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage;
                                                                                                                                                                                                                                                                                        Couture L, Stinchcomb D, Mcswiggen J, Bisgaier C,
                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC
(WARN ) WARNER LAMBERT CO.
                                                                                                                                                                                                                                                                                                                   WPI; 1996-321852/32.
                                                                                                                                                                                             11-DEC-1995;
                                                                                                             Homo sapiens
                                                                                                                                      WO9620279-A1
                                                                                                                                                                                                                       23-DEC-1994;
                                                                                                                                                                  04-JUL-1996.
                                                                                  LDL, ss.
  The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (i) at least 5 ribose residues 5 (i) at 2-C-allyl modification at position 4 of the ENA, (iii) at least to 1 i a 2-C-allyl modifications; and (iv) a 3-ead modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloaning of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                    Enzymatic nucleic acid molecules having a hammer-head motif - used for
the treatment of arthritis, induction of graft tolerance or treatment of
auto-immune diseases.
                                                                                                                                                                                                                                                                                           C, Draper K, Pavco P;
Wincott F, Matulic-Adamic J;
Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.4; DB 1; Length 15;
Pred. No. 2.98+02;
4; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 2 A; 3 C; 6 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                            Jarvis T,
                                                                                                                                                                                                                                                                                                        Usman N,
Modak A,
                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10; Page 166; 307pp; English.
                                                                                             94US-00354920.
94US-00363253.
94US-00393254.
95US-0039620.
95US-0043614.
95US-00043614.
95US-0000951P.
                                                                                                                                                                                                                                                                                         Stinchcomb DT,
Gustofson J, Us
Thompson JD, Mc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%;
Similarity 64.3%;
9; Conservative
                                                                       95WO-US015516.
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95US-00541365
                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-300653/30.
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention
                                                                                                                                                                                                                                                                                         Beigelman L,
Mcswiggen J,
Karpeisky A,
                                                                                                                                                    20-APR-1995;
02-MAY-1995;
04-MAY-1995;
                   WO9618736-A2
                                                                                                13-DEC-1994;
23-DEC-1994;
23-DEC-1994;
17-FEB-1995;
                                                                      22-NOV-1995;
                                                                                                                                                                                                                                    05-OCT-1995
                                                                                                                                                                                                           07-JUL-1995
                                           20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                             Enzymatic
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95WO-US016000. 94US-00363240.

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AAT49608-T49863 represent target sequences for the human cholesterol ester transfer protein (CETP) hammerhead (HH) ribozymes (see AAT49881-75017). CETP is a 74 kD glycoprotein that facilitates neutral lipid transfer between plasma lipoproteins. The numbering of the targets refers to the position of the cleavage site in full length CETP. The ribozyme binds to 5 nucleotides either side of this site, provided the Sequence UH is immediately upstream. The ribozymes are able to cleave mRNA from the gene encoding CETP, thereby blocking synthesis and/or expression of the mRNA. By inhibiting CETP, the reverse cholesterol transport (RCT) pathway can be inhibited (or eliminated) thereby (HDL), prolonging HDL half life, and therefore increasing HDL levels. The ribozymes can be used to treat conditions associated with abnormal levels of CETP, specifically familial hypercholesterolaemia, atherosclerosis, peripheral vascular disease, hypercholesterolaemia, hypoalphalipoproteinaemia, dyslipidaemia, castenosis. By inhibiting CETP, the levels of HDL and low angioplastic restenosis. By inhibiting CETP, the levels of HDL and low chenty lipoproteins (LDP), and the HDL:LDL ratio are favourably altered density lipoproteins (LDP), and the HDL:LDL ratio are favourably altered
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (a decrease in LDL levels, and a corresponding increase in HDL levels). The HH ribozymes can also be used diagnostically to study genetic drift and mutations in diseased cells, and to detect CETP mRNA. As the HH ribozymes target specific regions of the CETP gene, they have low non-
New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA -
useful for preventing or treating initial development, progression or
regression of vascular diseases, esp. familial hypercholesterolaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 85.7%; Pred. No. 2.9e+02;
Matches 12; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 BP; 5 A; 3 C; 5 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                          Claim 4; Page 30; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                175 ACGAGTCCAAGGCA 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT49707 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACGAGUUCAAGGCA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specific activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT49707;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT49707
ID AAT4
XX
AC AAT4
XX
DT 02-M
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Gaps

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Human CETP HH ribozyme target sequence #1056.

(first entry)

02-MAR-1997

SEX BX BX BX

AAT49705;

AAT49705 standard; RNA; 15 BP

RESULT 595 AAT49705

398 GAAGGICTICTACG 411

|| ||:|:|:||| GAGGGUCUUCUACG 15

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AAV66430;

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Hammerhead ribozyme; cholesterol ester transfer protein; mRNA cleavage; neutral lipid transfer; plasma lipoprotein; atherosolerosis; atherectomy; reverse cholesterol transport; high density lipoprotein; therapy; CETP; familial hypercholesterolaemia; dyslipidaemia; hypoalphalipoproteinaemia; peripheral vascular disease; hyperbetalipoproteinaemia; RCT; inhibitor; angioplastic restenosis; low density lipoprotein; diabetes; HDL; human;
       Human CETP HH ribozyme target sequence #1057.
                                                                                                                                                             94US-00363240.
                                                                                                                                            95WO-US016000.
                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC (WARN ) WARNER LAMBERT CO.
                                                                                           Homo Bapiens
                                                                                                            WO9620279-A1
                                                                                                                                            .1-DEC-1995;
                                                                                                                                                             23-DEC-1994;
                                                                                                                           14-JUL-1996
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AAT 49608-T49863 represent target sequences for the human cholesterol ester transfer protein (CETP) hammerhead (HH) ribozymes (see AAT49881-51017). CETP is a 74 kD glycoprotein that facilitates neutral lipid transfer between plasma lipoproteins. The numbering of the targets refers to the position of the cleavage site in full length CETP. The ribozyme binds to 5 nucleotides either side of this site, provided the sequence UH is immediately upstream. The ribozymes are able to cleave mRNA from the gene encoding CETP, thereby blocking synthesis and/or expression of the synthhiting CETP, the reverse cholesterol transport (RCT) pathway can be inhibited (or eliminated) thereby preventing the reduction in size density of the high density lipoproteins (HDL), prolonging HDL half life, and therefore increasing HDL levels of CETP, specifically familial conditions associated with abnormal levels of CETP, specifically familial hypercholesterolaemia, atherosolerosis, peripheral vascular disease, hyperbetalipoproteins disease, transplant, atherectony and canging laterase in the HDL:LDL ratio are favourably altered angioplastic restenosis. By inhibiting CETP, the levels of HDL levels). The HH ribozymes can also be used diagnostically to study genetic drift and mutations in diseased cells, and to detect CETP mRNA, As the HH crears of the carbozymes target specific regions of the CETP gene, they have low non-specific and mutations to the CETP gene, they have low non-specific and contents. H New ribozyme(s) for cleaving cholesterol ester transfer protein mRNA useful for preventing or treating initial development, progression or regression of vascular diseases, esp. familial hypercholesterolaemia. Pape M; Bisgaier C, Couture L, Stinchcomb D, Mcswiggen J, Claim 4; Page 30; 72pp; English. WPI; 1996-321852/32. specific activity

Seguence 15 BP; 5 A; 3 C; 5 G; 0 T; 2 U; 0 Other;

Gaps ö ch 2.9%; Score 12.4; DB 1; Length 15; 1 Similarity 85.7%; Pred. No. 2.9e+02; 12; Conservative 1; Mismatches 1; Indels Local Similarity Query Match Matches

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AAV66430 standard; DNA; 15 RESULT 597 AAV66430 ID AAV6 XX

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functioning as promoters for the tetracycline resistance (TetR) gene.

functioning as promoters for the tetracycline resistance (TetR) gene.

They are derived from the -35 promoter sequence of the TetR gene of

plasmid pBR322. The sequences were produced to exemplify the invention.

The specification describes a method for obtaining an oligonucleotide

that confers a predetermined biological function, such as regulation of

expression or a biological activity of a polypeptide, on a cell. The

method comprises cloning a heterogeneous pool of oligonucleotides into an

expression vector, where the clones oligonucleotides are transcribed or

cat as regulatory sequences, introducing a random sample of the cloned

oligonucleotides into a population of cells that do not exhibit the

predetermined biological function, selecting a subpopulation of cells

exhibiting the predetermined biological function from the selected

subpopulation of cells. The process is used, for example, for identifying

new forms of the Escherichia coli tetracycline resistance gene promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identification of biologically active DNA sequences - by transforming cells with random oligo-nucleotide (s).
                                                                                                 -35 promoter; plasmid pBR322; tetracycline resistance gene; TetR; promoter; Escherichia coli; active site; beta-lactamase gene; ss.
                                                               -35 promoter sequence of TetR gene of plasmid pBA6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Fig 3; 24pp; English.
                                                                                                                                                                                                                                                                                                                  89US-00368674.
92US-00881607.
93US-00105108.
                                                                                                                                                                                                                                                                                                   86US-00887070
                                                                                                                                                                                                                                                               94US-00316415
                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                         UNIW ) UNIV WASHINGTON.
                                                                                                                                                                                                                                                                                                                                                                                                                           Loeb LA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-582545/49.
                                                                                                                                                                                                                                                               30-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                         11-AUG-1993;
                                   06-JAN-1999
                                                                     Substituted
                                                                                                                                                                                            US5824469-A.
                                                                                                                                                                                                                               20-OCT-1998.
                                                                                                                                                                                                                                                                                                                    19-JUN-1989
                                                                                                                                                                                                                                                                                                                                                                                                                           Horwitz MS,
                                                                                                                                                          Synthetic
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Gaps ; 0 2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; ive 0; Mismatches 1; Indel8 92.98; Best Local Similarity 92.9 Matches 13; Conservative Query Match

Sequence 15 BP; 0 A; 5 C; 9 G; 1 T; 0 U; 0 Other; and the active site of the beta-lactamase gene

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AAC73241 standard; DNA; 15 (first entry) 02-FEB-2001 AAC73241; \$\$\$\$\$\$\$\$\$\$

뗦

RESULT 598

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Forward primer #43 used in multiplexing PCR/SBE assay.

Oligonucleotide array; genotyping; single base extension reaction; SBE;

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The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide polymorphism (SNP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                        Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                                                                                                                    Kaplan P, Lander ES, Lockhart DJ;
PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 15 BP; 4 A; 5 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                               (WHED ) WHITEHEAD INST BIOMEDICAL RES (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; Page 52; 70pp; English
                                                                                                                                                                                                                                                                                    Huang X,
                                                                                                                                                                           99US-0126473P.
99US-0140359P.
                                                                                                                                         27-MAR-2000; 2000WO-US008069.
                                                                                                                                                                                                                                                                                    Hirschhorn JN,
                                                                                                                                                                                                                                                                                                                                      WPI; 2000-656171/63.
                                                                                                                                                                                                                                                                                    Fan J, Hirschhorn
Ryder T, Sklar P;
                                                                    WO200058516-A2
                                                                                                                                                                                              23-JUN-1999;
                                    Unidentified
                                                                                                                                                                               26-MAR-1999;
                                                                                                        05-OCT-2000
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Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or

Example 8; Page 62; 201pp; English

inflammation.

Edmondson SR;

Werther GA,

Wraight CJ,

WPI; 2001-041421/05

(MURD-) MURDOCH CHILDRENS RES INST.

99US-0140345P

21-JUN-1999;

21-JUN-2000; 2000WO-AU000693

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Gaps
                                     .
0
Score 12.4; DB 1; Length 15; Pred. No. 2.9e+02; 0; Mismatches 1; Indels
     Query Match
Best Local Similarity 92.9%;
Matches 13; Conservative
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388 ACGGCGCCAAGAAG 401 1 ACGGCGCCAAGATG 14 ö 임

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Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; Keloid; skin discorder; Insulin-like Growth Factor. I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keardosls; neophasia; gcleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplama; kidney disease; neoblation of the retina; ss.
AAF49243 standard; DNA; 15 BP
                                                                                                        IGF-I oligonucleotide #203.
                                                                      (first entry)
                                                                      30-MAR-2001
                                   AAF49243;
BXBXSXXXXXXXXXXXXXXXXXXX
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21-JUN-2000; 2000WO-AU000693

WO200078341-A1

28-DEC-2000

Homo sapiens

WO200078341-A1.

28-DEC-2000.

Homo sapiens

99US-0140345P.

21-JUN-1999;

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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleoride, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, cinflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide which can be useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ubb, pilaris, serborrhoea, keloids, keratosis, ichthyosis, pityriasis, ubt, pilaris, serborrhoea, keloids, keratosis, chthyosis, pityriasis, warts, benign growths, cancers of the skin, a hyperneovascular condition out the retina, brain or skin, growth factor-mediated malignancies, other sclerotic brain or skin, growth factor-mediated malignancies, other sclerotic brain or standard and propresent of the inside of blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; sein discorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFB-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match

2.9%; Score 12.4; DB 1; Length 15.

Best Local Similarity 92.9%; Pred. No. 2.9e+02;

Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 2 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF53588 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IGF-I oligonucleotide #4548.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 GGTGACCGAGGGCT 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGTGATCGAGGGCT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF53588;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ઠે
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The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligomucleotide, (for Insulin-like Growth Factor [167]-1 receptor, 1GF binding protein [1678]-2 or 1GFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, configuration and/or other disorders. The present sequence is an oligomucleotide which can be used to design the antisense oligomucleotide which can be used to design the antisense oligomucleotide which can be useful for ameliorating the effects of psoriasis, chthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, chrain or skin, growth factor—mediated malignancies, other sclerotic disease, kidney disease, hyperplasia
                                                                                                                                                           Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 3 A; 1 C; 7 G; 4 T; 0 U; 0 Other;
                                                                  Werther GA, Edmondson SR;
                                                                                                                                                                                                                                                                                Example 8; Page 90; 201pp; English.
                 (MURD-) MURDOCH CHILDRENS RES INST
                                                                                                               WPI; 2001-041421/05.
                                                                                                                                                                                                                                    inflammation.
                                                                  Wraight CJ,
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1; Indels 0; Gaps Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indele 356 CAGCGACTTCCTCA 369 15 cádccácrrcíca 2 ठ 8

AAF53590 standard; DNA; 15 BP. IGF-1 oligonucleotide #4550. 30-MAR-2001 (first entry) AAF53590; RESULT 601 AAF53590, 

Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding procein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;

Homo sapiens.

WO200078341-A1 28-DEC-2000 21-JUN-2000; 2000WO-AU000693.

99US-0140345P. 21-JUN-1999; (MURD-) MURDOCH CHILDRENS RES INST.

Werther GA, Edmondson SR; Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 90; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an anticomplete clede. (for Insulin-like Growth Factor IIGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation.

Inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the artisense oligonucleotide which can be used to design the artisense oligonucleotide which can be used to design the artisense oligonucleotide is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor—mediated malignancies other sclerotic disease, kidhey disease, hyperpoliferation of the inside of blood

Sequence 15 BP; 3 A; 2 C; 7 G; 3 T; 0 U; 0 Other;

0; Gaps 2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; ative 0; Mismatches 1; Indels 2.9% Best Local Similarity 92.9% Matches 13, Conservative

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AAF49244 standard; DNA; 15 BP. RESULT 602 AAF49244

IGF-I oligonucleotide #204. (first entry) 30-MAR-2001 

AAF49244;

Antiëense therapy, antiproliferative, antiinflammatory, antipeoriatic, cytostatic, dermatological; cardiant; virucide, ophthalmological; keloid; skin discorder, insulin-like Growth Factor. I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scaleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition; hyperplasia; kidney disease;

Homo sapiens.

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693

(MURD-) MURDOCH CHILDRENS RES INST.

21-JUN-1999; 99US-0140345P.

Edmondson SR, Werther GA, Wraight CJ,

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering

UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 62; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [10F]-1 receptor, IGF binding protein [10FBP]-2 or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inhibiting or reducing growth factor mediated cell proliferation, inhibiting or reducing growth factor mediated cell proliferation, oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliotating the effects of psoriasis, ichthyosis, pityriasis, ruba, pliaris, serborrhoea, keloids, keratosis, ichthyosis, pityriasis, ruba, pliaris, serborrhoea, keloids, keratosis, hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, thing the proposed of the inside of blood vessels or any other hyperplasia

Sequence 15 BP; 3 A; 2 C; 7 G; 3 T; 0 U; 0 Other;

1; Indels 0; Gaps Ouery Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels

20 GGTGACCGAGGGCT 33

δ

RESULT 603 AAF49333

AAF49333 standard; DNA; 15 BP. 

AAF49333;

30-MAR-2001 (first entry)

IGF-I oligonucleotide #293.

Antisense therapy, antiproliferative, antinflammatory, antipsoriatic, cytostatic; dermatological, cardiant, virucide, ophthalmological, keloid, skin disorder, Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding procein, IGFBP-2; IGFBP3; inflammation, psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neophasia; scleroderma; wart; skin cancer; sclerotic disease; hypermeowascular condition, hyperplasis; kidney disease; necovascular condition of the retina; ss.

Homo sapiens.

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693.

99US-0140345P. 21-JUN-1999;

Wraight CJ, Werther GA, Edmondson SR; (MURD-) MURDOCH CHILDRENS RES INST.

WPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

The present invention relates to a method for ameliorating the effects of artisense oligonuclecide, (for Insulin-like Growth Factor [IGP]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present egquence is an oligonuclectide which can be used to design the antisense oligonuclectides of the present invention (see AAF45151 and AAF45153-F4561). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, hepplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovacular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood c hyperp...
A; 3 C; 4 G; 6 T; 0 U; 0 Other,
2.9%; Score 12 4; DB 1; Length 15;
y 92.9%; Pred, No. 2.9e+02;
y 92.9%; Pred, Numatches 1; Indels 0; Gaps Sequence 15 BP; 2 A; 3 C; 4 G; 6 T; 0 U; 0 Other; Example 8; Page 62; 201pp; English. vessels or any other hyperplasia 112 ACCGCAGCAAGTAC 125 15 ACAGCAGCAAGTAC 2 Local Similarity 92.9 les 13; Conservative Query Match Matches ઠે

AAF49334 standard; DNA; 15 BP. RESULT 604 AAF49334/c

Antisense therapy, antiproliferative, antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; insulin-like Growth Pactor 1 receptor; 1GF-1; pityriasis; IGF binding protein; IGFBP-3; inflammation; psoriasis; pitaris; growth factor mediated cell proliferation; ichthyosis; serborrhoes; ruba; keratosis, mediated cell proliferation; ichthyosis; serborrhoes; ruba; hyperneovascular condition, hyperplasis; kinney disease; neovascular condition of the retina; ss. 

IGF-I oligonucleotide #294. 30-MAR-2001 (first entry)

AAF49334;

WO200078341-A1.

28-DEC-2000.

21-JUN-2000; 2000WO-AU000693. 99US-0140345P. 21-JUN-1999; Wraight CJ, Werther GA, Edmondson SR;

(MURD-) MURDOCH CHILDRENS RES INST.

VPI; 2001-041421/05.

Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.

Example 8; Page 62; 201pp; English.

The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an

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The invention relates to an isolated polynucleotide comprising a sequence which is a polymorphic variant of a reference sequence for crystallin, where the polymorphic variant comprises a CRYBB1 isogene defined by a haplotype from haplotypes 1-16 as given in the specification. Also included are a transgenic non-human animal transformed or transfected with the polymorphic variant, a computer system for storing and analysing polymorphism data for CRYBB1 isogenes, methods of determining an individuals haplotype or genotype as well as methods of determining the association of a particular haplotype with a disease or trait and a secondosition comprising at least one genotyping oligonucleotide
                receptor, 10F binding protein [16FBP] or 1GFBP3, which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense passent invention (see AAF45151 and AAF45153 - F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, planis, serbornhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperpoliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, crystallin beta B1, CRYBB1; chromosome 22q12.1; ophthalmalogical, cataract, allele specific oligonucleotide, ASO, ss; haplotype; genotyping; transgenic animal; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                       1; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel polymorphic variants of crystallin, beta B1 useful in studying expression and function of the protein, useful for screening candidate drugs to treat diseases e.g. cataract.
antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1
                                                                                                                                                                                                                                                                                                                                          2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred, No. 2.9e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Choi JY, Kazemi A, Kliem SE, Koshy B, Rounds E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer #1 for human CRYBB1 gene haplotype PS10.
                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 1 A; 3 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 28; Page 31; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS97386 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-MAY-2000; 2000US-0202253P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-MAY-2001; 2001WO-US014715.
                                                                                                                                                                                                                                                                                                                                                                                                                              112 ACCCCAGCAGTAC 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14 ACAGCAGCAAGTAC 1
                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 92.9°
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-062253/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS97386;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 605
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(especially allele-specific oligonucleotides (ASO)) for detecting a polymorphism in the CRYBB1. The isogenes or haplotypes are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with CRYBB1 activity, e.g. cataract. and can also be used by the pharmaceutical research scientist to validate CRYBB1 as candidate target for, and in design of clinical trials of candidate drugs for, irreating a specific condition drugs or disease predicted to be associated with CRYBB1 activity. The ASOs are useful as probes and primers, and for assaying a polymorphism in the target region. The present sequence is a pck primer which amplifies a region of CRYBB1 containing one of 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting a base at a pre-determined position in a nucleic acid molecule, comprises performing primer extension reactions using base-specific detection primers in the presence of a nucleotide-degrading enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for detecting a base at a predetermined position in a nucleic acid molecule. The method comprises performing primer extension reactions using base-specific detection primers, each being specific for a particular base at the predetermined position. The allele-specific (AS) primer extension assay method of the invention is useful for detecting an allele-specific base at a predetermined position in a nucleic acid molecule, for high throughput single nucleotide polymorphism (SNP) analysis, and for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, allele-specific base detection; primer extension reaction; base-specific detection primer, allele-specific primer extension assay; AS; high throughput; single mucleotide polymorphism; SNP analysis; mutation detection; genetic variation; allele-specific extension; probe;
                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                       y Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 92.9%; Pred. No. 2.9e+02; nes 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                    Sequence 15 BP; 3 A; 4 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genomic DNA p53 SNP AS extension probe #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lundeberg J, Ahmadian A, Nyren P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 33; 59pp; English.
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23-FEB-2001; 2001US-00791190.
07-FEB-2002; 2002US-00071926.
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                                                                                                                                                                                                                                                                                                                                                                                                                132 CTGGCCCGCCTGGC 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                            polymorphic sites
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Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

exemplification of the invention

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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonuclectide having a first and second strand with at least a region of complementarity in between them. The first oligonuclectide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonuclectides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequence is an annealing oligonuclectide for Kan- target. This sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of
mutations and genetic variations. The new method solves the deficiencies of previous methods by providing a method of allele-specific extension that allows accurate discrimination between marched and mismarched configurations, as well as reducing or eliminating false positive results observed in prior art. The use of two allele-specific primers increases the sensitivity by a factor of two because signals of two extensions are obtained. The present sequence represents a probe used in the examples of
                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                             2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; iive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detection; purification; double D-loop formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UDS15G annealing oligonucleotide for Kan- target.
                                                                                                                                                            Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Usher MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                                                                                                      238 GAGGCTGCTTCCCG 251
                                                                                                                                                                                                                                                                                                                                                                                                 AAD48683 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mod base=
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                                                                                                                                the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-046824/04.
                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200279495-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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                                                                                                                                                                                                    Query Match
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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonucleotide having a first and second strand with at least a region of complementarity in between them. The first oligonucleotide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not obound. The methods, oligonucleotides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is an oligonucleotide which is used for determination of optimal oligonucleotide composition for the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonuclectide having a first and second strands with a region of
                             Gaps
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2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                               *tag= a
'note= "Locked nucleic acid (LNA)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= c
/note= "Locked nucleic acid (LNA)"
                                                                                                                                                                                                                                                                           Detection; purification; double D-loop formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Usher MG;
                                                                                                                                                                                                                                            Oligo O used for double D-loop formation.
                                                                                                                                                                                                                                                                                                                                    cocation/Qualifiers
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/note= "DNA"
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28-SEP-2001; 2001US-0325828P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complementarity in between.
                                                             155 CGGCTTCGACTGGG 168
                                                                                                                                                     1672/c
AAD48672 standard; DNA; 15
                                                                                                                                                                                                                      (first entry)
                                                                                     1s cadetracaacteds 2
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                Local Similarity 92.9 (es 13; Conservative
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*tag=
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                                                                                                                                                                                                                                                                                                          Unidentified
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         Query Match
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Page 293

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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonuclectide having a first and second strand with at least a region of complementarity in between them. The first oligonuclectide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonuclectides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is a locked nucleic acid (INA) which is used as an annealing oligonuclectide for Kan- target. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonuclectide having a first and second strands with a region of complementarity in between.
                                                                                                                                                                                    Detection; purification; double D-loop formation; locked nucleic acid;
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2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                   KLO2 annealing oligonucleotide for Kan- target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Usher MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 11; Page 49; 99pp; English.
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28-SEP-2001; 2001US-0325828P.
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                AAD48685 standard; DNA; 15
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                                                                                                      (first entry)
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                                                             AAD48685;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               double D loop at a target sequence within a double-stranded nucleic acid.

The method involves contacting the double-stranded nucleic acid with an oligonucleotide having a first and second strand with at least a region of complementarity in between them. The first oligonucleotide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonucleotides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic generate neomycin phosphorransferase mutant (Kan-) gene. This sequence is used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a novel method of producing a stabilised double D loop at a target semience within a double D loop at a target semience within a double.
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                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      double D-loop formation; neomycin phosphotransferase
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                                                                                                                                                                                                                                                                                                                                                                                                                             Oligo KLO2 used to generate neomycin phosphotransferase mutant
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                                   / Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 92.9%; Pred. No. 2.9e+02; les 13; Conservative 0; Mismatches 1; Indels
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Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
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28-SEP-2001; 2001US-0325828P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-MAR-2002; 2002WO-US009691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           complementarity in between.
                                                                                                                                  155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                          AAD48681 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cecciaceaciese 2
                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                 15 CGGCTACGACTGGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-046824/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           purification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   40200279495-A2.
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Inidentified.

Detection;

10-OCT-2002

Kmiec EB,

15

g

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24-FEB-2003

AAD48681;

RESULT 609

AAD48681

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Gaps

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Wed Apr 21

10-OCT-2002

Kmiec EB,

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The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonucleotide having a first and second strand with at least a region of complementarity in between them. The first oligonucleotide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonucleotides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is an oligonucleotide which is used for double D-loop formation. This sequence is used in the
                                                                                                                                                                                                                                 Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonuclectide having a first and second strands with a region of complementarity in between.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid, hammerthead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; anti-HCV; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-HCV enzymatic nucleic acid substrate sequence #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 BP, 2 A, 6 C, 5 G, 2 T, 0 U, 0 Other;
                                                                                                                                          Usher MG;
                                                                                                                                                                                                                                                                                                                                                Example 2; Page 35; 99pp; English.
                                                                                                                                        Rice MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACD66419 standard; RNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-2001; 2001US-00817879.
08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-03370559P.
                      27-MAR-2001; 2001US-0279146P.
28-SEP-2001; 2001US-0325828P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155 CGGCTTCGACTGGG 168
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                                                                                         (UYDE ) UNIV DELAWARE.
                                                                                                                                          Gamper HB,
                                                                                                                                                                                       WPI; 2003-046824/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200281494-A1
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08-JUN-2001;
24-OCT-2001;
05-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-0CT-2002
                                                                                                                                        Kmiec EB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           à
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a novel method of producing a stabilised double D loop at a target sequence within a double-stranded nucleic acid. The method involves contacting the double-stranded nucleic acid with an oligonuclectide having a first and second strand with at least a region of complementarity in between them. The first oligonuclectide strand has a region that is complementary to a first strand of the target and binds to the recombinase while the second strand is not bound. The methods, oligonuclectides, compositions and kits are useful for detecting and purifying known nucleic acid targets and for manipulating defined nucleic acid target sequences. The present sequence is an annealing oligonuclectide for Kan- target. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Producing a stabilized double D loop at a target sequence within a double stranded nucleic acid comprises contacting the nucleic acid with an oligonucleotide having a first and second strands with a region of complementarity in between.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detection; purification; double D-loop formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 2 A; 6 C; 5 G; 0 T; 2 U; 0 Other;
                                          Location/Qualifiers
1. 15
7-tag= 7
7-dd base= OTHER
7-note= "2'-0-methyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Usher MG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligo N used for double D-loop formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 11; Page 48; 99pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Rice MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 612
AAD48648/c
ID AAD48648 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-MAR-2002; 2002WO-US009691
                                                                                                                                                                                                                                                                                27-MAR-2002; 2002WO-US009691
                                                                                                                                                                                                                                                                                                                             27-MAR-2001; 2001US-0279146P.
28-SEP-2001; 2001US-0325828P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-FEB-2003 (first entry)
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gamper HB,
                                                                                                                                                                                                                                                                                                                                                                                                (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-046824/04.
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                                                                                                                                                                                         WO200279495-A2
                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified.
Unidentified
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AAD48648;

#X2XEXEXEXEXEX

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Gaps ö 08-JUN-2001; 2001US-00877478. 08-JUN-2001; 2001US-0296876P. 08-JUN-2001; 2001US-0335059P. 05-DEC-2001; 2001US-0337055P.

RIBOZYME PHARM INC

RIBO-)

(BLAT/) BLATT L.
(MACE/) MACKEAK D.
(MCSW/) MCSWIGGEN J.
(MORK/) MORRISSEY D.
(PAVC/) PAVCO P.
(LREP/) LEE P.
(LREP/) LEE P.
(ROBY/) DRAFER K.
(ROBZ/) ROBERTS E.

2001US-00817879

26-MAR-2002; 2002WO-US009187

WO200281494-A1 17-0CT-2002

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptuse primer sequences, as well as oligonucleotides that specifically bind the Bnhancer I region of HBV genes and dor HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the carcinoma. The present sequence represents a substrate for one of the carcinoma. The present sequence acid sequences disclosed in the present
                                                                                                                                                                                                                                                                             Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                           Mcswiggen J, Morrissey D, Pavco P, Lee P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 2 A; 6 C; 2 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 326; 387pp; English
                  RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 92.5.
Best Local 3; Conservative
                                                                                                                                                                                                            Roberts E;
                                   BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                           Macejak D,
                                                                                                                                                                                                                                            WPI; 2003-229207/22
                                                                                                                      LEE P.
DRAPER K.
                                                                                                                                                          ROBERTS E.
                                                                                                                                                                         Blatt L, Me
                                                                                                                                                                                                                                                                                                                    nfection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nvention
                                                                                                    (PAVC/)
(LEEP/)
(DRAP/)
(ROBE/)
                                 (BLAT/)
(MACE/)
(MCSW/)
(MORR/)
```

Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus

Mcswiggen J, Morrissey D, Pavco P, Lee P;

Macejak D, Roberts E;

Blatt L, Me Draper K, I

WPI; 2003-229207/22

infection.

```
The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HRV) RNY. The nucleic acid molecules include antisense and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, an eargmatic nucleic acids such as harmerhead ribozymes, DNAzymes, and enzymatic nucleic acid decoy molecules and G-cleaver ribozymes, Also disclosed are uncleic acid decoy molecules and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and chose the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and generations are useful. For the treatment of degenerative and captement as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represence a target for one of the anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Microarray; probe; Mycobacterium; antibiotic-resistance; genotyping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 2 A; 6 C; 2 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mycobacterium gastrii specific probe GAS-03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 322; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACC73353 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 GGAGTGAAACTGCG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 GGAGTGAAAATGCG 2
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ID ACC7
XX
AC ACC7
XX
DT 15-J
XX
KW Micr
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0; Gaps

2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; Artive 0; Mismatches 1; Indels

6 GGAGTGAAACTGCG 19

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15 GGAGTGAAAATGCG 2

Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; harmerihead ribozyme; DNAzyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; anti-HCV; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antihifammatory; target; ss.

Hepatitis C virus

Anti-HCV nucleic acid molecule target sequence #232.

23-SEP-2003 (first entry)

ACD66349;

ACD66349 standard; RNA; 15 BP.

RESULT 614 ACD66349/c

17-APR-2003

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The present sequence is that of kan klo3, an oligonucleotide mismatched (non-hybridising) to the triplet repeat region of exon 1 of the human Huntington's disease (HD) gene. The oligonucleotide is modified by including locked nucleic acid (InA) residues at both ends. Administration of this short, modified oligonucleotide to neuronal PC12 cells bearing an HD exon 1-GFP fusion gene did not result in a decrease in Huntington protein (huntingtin) aggregation in cell culture studies. The invention realases to oligonucleotides, including oligonucleotides containing IMA modifications, that alter the genomic HD gene sequence and/or reduce the propensity of huntingtin to form intracellular aggregates. These can be used for the treatment or prevention of HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
'note= "locked nucleic acid"
                                                                       note= "locked nucleic acid"
                                                                                                           *tag= e
mod_base= OTHER
'note= "locked nucleic acid"
                                                                                                                                                                                  *tag= f
/mod_base= OTHER
/note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                      /note= "locked nucleic acid"
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                                                                                                                                                                                                                                                                   /*tag= g
/mod_base= OTHER
                                    '*tag= d
'mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-AUG-2001; 2001US-0310757P.
08-AUG-2001; 2001US-0310770P.
04-AUG-2001; 2001US-0337219P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kmiec EB, Parekh-Olmedo H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15 cecraceacrees
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                                                                                                                                                                        modified base
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                      modified base
                                                                                               modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a microarray comprising a support, a first probe for genotyping Mycobacterium species, second probe for differentiating Mycobacterium thebroulosis strains, and a third probe for detecting antibiotic-resistant strains, where the probes are immobilized on the support. This sequence represents an example of the first probe used for genotyping Mycobacterium species. The array is useful for simultaneously genotyping Mycobacterium species, differentiating M. tuberculosis strains and detecting antibiotic-resistant strains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                   Microarray for simultanecusly genotyping Mycobacteria species, differentiating Mycobacterium tuberculosis strains and detecting antibiotic-resistant strains, comprises specific probes on a support.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Huntington's disease; nootropic; anticonvulsant; huntingtin; human; locked nucleic acid; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . Match 2.9%; Score 12.4; DB 1; Length 15; Local Similarity 92.9%; Pred. No. 2.9e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Locked nucleic acid-containing oligonucleotide kan klo3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15 BP; 1 A; 3 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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                                                                                                                                                                                                                                                                                            Song E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Page 57; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
                                                                                                                                                                              09-OCT-2001; 2001KR-00062125
                                                                                                                                        09-OCT-2002; 2002WO-KR001885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             175 ACGAGTCCAAGGCA 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ81751 standard; DNA; 15
                                                                                                                                                                                                                   SJ HIGHTECH CO LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 ACGAGICCAAGACA 2
                                                                                                                                                                                                                                                                                            Kim C, Park H, Jang H,
                            Mycobacterium gastrii
                                                                                                                                                                                                                                                                                                                                  WPI; 2003-403109/38.
                                                                                                                                                                                                                   (SJHI-) SJ HIGHTI
(KIMC/) KIM C.
(PARK/) PARK H.
                                                                WO2003031654-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
modified_base
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Query Match Best Loca Matches

8

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Gaps

ö

Synthetic

ABZ81751;

RESULT 616 ABZ81751 07-AUG-2002; 2002WO-US025352

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Huntington's disease; nootropic; anticonvulsant; huntingtin; human; locked nucleic acid; gene therapy; ss.
Locked nucleic acid-containing oligonucleotide kan klo2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= o
/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic acid"
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mod_base= OTHER
'note= "locked nucleic acid"
                                                                                                                                                                                                                              *tag= b
/mod_base= OTHER
/note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag= e
mod_base= OTHER
fnote= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag= f
mod_base= OTHER
note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            *tag= i
mod_base= OTHER
note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                  *tag= c
mod_base= OTHER
note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                      *tag= d
|mod_base= OTHER
|note= "locked nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= g
/mod_base= OTHER
/note= "locked nucleic acid"
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                                                                                                                                                                                                   note= "locked nucleic acid"
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/mod_base= OTHER
/note= "locked nucleic
                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= h
/mod_base= OTHER
                                                                                                                                                                  *tag= a
'mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003013437-A2
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                         modified base
                                                                                                                                                                                                                                                                                    modified base
                                                                                   Homo sapiens
                                                                                                   Synthetic.
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20-FEB-2003

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                                                                                                                                                                                                                                                                                                                The present sequence is that of kan klo1, an oligonuclectide mismatched (non-hybridising) to the triplet repeat region of exon 1 of the human lumining on's disease (HD) gene. The oligonuclectide is modified by having locked nucleic acid (LNA) residues throughout its length. Administration of this short, modified oligonuclectide to neuronal PC12 cells bearing an protein (huntingtin) aggregation in cell culture studies. The invention relates to oligonuclectides, including oligonuclectides containing LNA modifications, that alter the genomic HD gene sequence and/or reduce the propensity of huntingtin to form intracellular aggregates. These can be used for the treatment or prevention of HD
                                                                                                                                                                                                         New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Huntington's disease gene non-specific oligonucleotide Kan uD7T/15G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Huntington's disease; nootropic; anticonvulsant; phosphorothioate; huntingtin; human; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
2.9%; Score 12.4; DB 1; Length 15;
Best Local Similarity 92.9%; Pred. No. 2.9e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
1. 15
/*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                         Example 5; Page 71; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ81742 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-AUG-2001; 2001US-0310757P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-AUG-2002; 2002WO-US025352.
                                      07-AUG-2001; 2001US-0310757P.
08-AUG-2001; 2001US-0310770P.
08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       155 CGGCTTCGACTGGG 168
                                                                                                                                                   Kmiec EB, Parekh-Olmedo H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 CGGCTACGACTGGG 2
                                                                                                                  (UYDE ) UNIV DELAWARE
                                                                                                                                                                               WPI; 2003-256478/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003013437-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ81742;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 618
ABZ81742/c
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The present sequence is that of single-stranded oligonucleotide Kan MuR/1SG, which has 2'0-Me modifications throughout its length.

Administration of this oligonucleotide to PCI2 neuronal cells containing an engineered Huntington's disease (HD) gene exon 1 including alternating, repeating Gln codons (CAA/G) had little effect on HD protein (huntingtin) aggregation. This was in contrast to other modified oligonucleotides (see ABZB173'-39) which, although non-specific and non-hybridising to the HD gene, and being incapable of directing sequence hybridising of the tripler repeat region of exon 1, nevertheless reduced the formation of HD protein aggregates. Such oligonucleotides can be used for the treatment or prevention of HD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to distinguishing presence of a nonsupercoiled target nonsupercoiled target
                                                                                      New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington a disease gene to be altered, useful for treating or preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nonsupercoiled nucleic acid; target query region; genotyping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.96+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 11; SEQ ID NO 42; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide of the invention #42.
                                                                                                                                                                            Example 1; Page 59; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC13797 standard; DNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-SEP-2001; 2001US-0325828P.
27-MAR-2002; 2002WO-US009691.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-SEP-2002; 2002WO-US031073.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155 CGGCTTCGACTGGG 168
                        Parekh-Olmedo H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 cecraceacrece 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYDE ) UNIV DELAWARE
                                                        WPI; 2003-256478/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-371937/35.
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                        Kmiec EB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kmiec EB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC13797;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC13797
       ਨੇ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                      The present sequence is that of single-stranded phosphorothioate oligonucleotide Kan ub77/15G. Administration of this oligonucleotide to oligonucleotide college containing an engineered Huntington's disease (HD) gene exon including alternating, repeating Gln codons (CAA/G) resulted in a reduction in the formation of HD protein (huntingtin). Kan ub77/15G is an example of modified oligonucleotides of the invention, which, although non-specific and non-hybridising to the HD gene, and incapable of directing sequence alternation of the triplet repeat region of exon 1, nevertheless reduce the formation of HD protein aggregates. Such oligonucleotides can be used for the treatment or prevention of HD
                                                                                                                                                                        New single stranded oligonucleotides comprising a DNA domain having at least one mismatch with respect to the genetic sequence of the Huntington's disease gene to be altered, useful for treating or preventing Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    duntington's disease; nootropic; anticonvulsant; huntingtin; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Huntington's disease gene non-specific oligonucleotide Kan uR/15G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "all RNA 2'-0-Methyl modifications"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                          Example 1; Page 60; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ81741 standard, RNA, 15 BP.
08-AUG-2001; 2001US-0310770P.
08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-AUG-2001; 2001US-0310757P.
08-AUG-2001; 2001US-031070F.
08-AUG-2001; 2001US-0310889P.
04-DEC-2001; 2001US-0337219P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-AUG-2002; 2002WO-US025352.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ...15
*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cuery Match
Best Local Similarity 92.9°
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15 céécraceacreée 2
                                                                                                   Kmiec EB, Parekh-Olmedo
                                                                   (UYDE ) UNIV DELAWARE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYDE ) UNIV DELAWARE
                                                                                                                                      WPI; 2003-256478/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003013437-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-FEB-2003
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Gaps

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target by a nucleotide within a common target query region (TQR), trayleting a nucleotide within a common target query region (TQR), trayleting using a recombinase to mediate formation of deproteinizationstable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target mucleic acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids and is also useful for sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acid target from there in onsupercoiled double- stranded nucleic acids within a separating a nonsupercoiled double- stranded nucleic acids within a sumple of nucleic acids within a sample of nucleic acids within a sample of nucleic acids within a large functed. The methods are readily multiplexed, permitting a large number of loci to be screened within a single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic sequence is an oligonucleotide of the invention. Offers
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Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

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0; Gaps
/ Match
2.9%; Score 12.4; DB 1; Length 15;
Local Similarity 92.9%; Pred. No. 2.9e+02;
les 13; Conservative 0; Mismatches 1; Indels
                                                 Matches
                                                                                           à
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ADC13793 standard; DNA; 15 BP. ADC13793; RESULT 621 ADC13793/c

Oligonucleotide of the invention #38. (first entry) 18-DEC-2003 

nonsupercoiled nucleic acid; target query region; genotyping; ss.

Synthetic.

WO2003027640-A2.

03-APR-2003.

27-SEP-2002; 2002WO-US031073. 28-SEP-2001; 2001US-0325828P. 27-MAR-2002; 2002WO-US009691.

(UYDE ) UNIV DELAWARE.

Kmiec EB,

WPI; 2003-371937/35

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from

Example 10; SEQ ID NO 38; 179pp; English.

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinization-

ctable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a monsupercolled target mucleic acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercolled target variants within a sample of mucleic acids and is also useful for targets within the sample of nucleic acids and is also useful for separating a nonsupercolled double- stranded nucleic acids within a sample of nucleic acids within a larget of nucleic acids within a sample of nucleic acids within a sample of succeed within a separating a large number of loci to be screened within a single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic selectivity, with or without contemporaneous detection, offers sequence is an oligonucleotide of the invention.

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Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;

ö Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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1760/c ADC13760 standard; DNA; 15 BP. RESULT 622 ADC13760/

18-DEC-2003 (first entry) ADC13760; 

nonsupercoiled nucleic acid; target query region; genotyping; ss.

Oligonucleotide of the invention #5.

Synthetic.

WO2003027640-A2.

03-APR-2003.

27-SEP-2002; 2002WO-US031073.

28-SEP-2001; 2001US-0325828P. 27-MAR-2002; 2002WO-US009691.

(UYDE ) UNIV DELAWARE.

Kmiec EB, Rice MC;

WPI; 2003-371937/35.

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from variant.

Example 2; SEQ ID NO 5; 179pp; English.

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinization-stable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic

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acid such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acid target from other nonsupercoiled double- stranded nucleic acid target from where 10-10000 fold purification is effected. The methods are readily where 10-10000 fold purification is effected. The methods are readily single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
             %$GGGGGGGGGGGG
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Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels 155 CGGCTTCGACTGGG 168 15 CGGCTACGACTGGG 2 ઠે g

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Gaps

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ADC13784 standard; DNA; 15 BP (first entry) 18-DEC-2003 ADC13784; RESULT 623 ADC13784/

nonsupercoiled nucleic acid; target query region; genotyping; ss. Oligonucleotide of the invention #29. 27-SEP-2002; 2002WO-US031073. WO2003027640-A2. 03-APR-2003 Synthetic. 

28-SEP-2001; 2001US-0325828P. 27-MAR-2002; 2002WO-US009691. (UYDE ) UNIV DELAWARE

Kmiec EB, Rice MC;

WPI; 2003-371937/35

Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target query region which distinguish target from

Example 11; SEQ ID NO 40; 179pp; English.

Example 5; SEQ ID NO 29; 179pp; English

The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinization-stable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic cacid such as a linear duplax DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled

The present invention relates to distinguishing presence of a monsupercoiled target nucleic acid from presence of nonsupercoiled target variants within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinizationstable double D loop in TQR and then distinguishing degree of formation of double D loops that are stable to deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic coil such as a linear duplax DWA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acid target from other nonsupercoiled double- stranded nucleic acid target from other nonsupercoiled double- stranded nucleic acids target from

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targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded mucleic acid target from cher nonsupercoiled mucleic acids within a sample of nucleic acids, where 10-10000 fold purification is effected. The methods are readily multiplexed, permitting a large number of loci to be screened within a single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Distinguishing nonsupercoiled target nucleic acid in sample of nucleic acids from variants of the target, by forming deproteinization-stable double D loops in target from
                                                                                                                                                                                                      Query Match 2.9%; Score 12.4; DB 1; Length 15; Best Local Similarity 92.9%; Pred. No. 2.9e+02; Matches 13; Conservative 0; Mismatches 1; Indels ; 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nonsupercoiled nucleic acid; target query region; genotyping; ss.
                                                                                                                                                                            Sequence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide of the invention #40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-SEP-2002; 2002WO-US031073.
                                                                                                                                                                                                                                                                155 CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                   ADC13795/c
ID ADC13795 standard; DNA; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                               ADC13795;
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Gaps

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The present invention relates to distinguishing presence of a nonsupercoiled target nucleic acid from presence of nonsupercoiled target nucleic acids the variants differing from target by a nucleotide within a sample of nucleic acids, the variants differing from target by a nucleotide within a common target query region (TQR), involving using a recombinase to mediate formation of deproteinization. It compared to be a common target guery region (TQR), involving using a recombinase to mediate formation of deproteinization. The method is useful for distinguishing the presence of a nonsupercoiled target nucleic acids such as a linear duplex DNA, covalently closed circle, or artificial chromosome from the presence of nonsupercoiled target variants within a sample of nucleic acids. The method distinguishes several nonsupercoiled targets within the sample of nucleic acids and is also useful for separating a nonsupercoiled double- stranded nucleic acid target from other nonsupercoiled nucleic acids within a sample of nucleic acids, where 10-1000 fold purification is effected. The methods are readily multiplexed, permitting a large number of loss screened within a construction and an arriver.
where 10-10000 fold purification is effected. The methods are readily multiplexed, permitting a large number of loci to be screened within a single sample, may be adapted to a variety of existing detection systems, and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single sample, may be adapted to a variety of existing detection systems
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nonsupercoiled nucleic acid; target query region; genotyping; ss.
                                                                                                                                                                                                                                         2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02;
                                                                                                                                                                                                Seguence 15 BP; 2 A; 6 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                     0; Mismatches
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27-MAR-2002; 2002WO-US009691.
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                                                                                                                                                                                                                                                                                                                              CGGCTTCGACTGGG 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC13796 standard; DNA; 15
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                                                                                                                                                                                                                                                                                     13; Conservative
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                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003027640-A2.
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The invention relates to a novel method for amplifying a DNA using polymerase chain reaction (PCR) comprising synthesising the first region of a base sequence to be amplified by designing a pair of primers so as to place the first region between them and to anneal each other at the 3-end and carrying out a polymerase chain reaction (PCR) using the primers. Subsequently, the second region is synthesised by designing a pair of primers so as to place the second region partly overlapping with the first region of the base sequence between them and to anneal each other at the 3'-end and carrying out a PCR using the primers. Pinally, the first region is annealed to the second region generating the template to carry out a PCR and thus to synthesize a base sequence containing the first and the second regions. The method of the invention may be useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amplification of a DNA, a gene encoding the repeated sequence of an amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence is that of the mucin-
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                                                                                                                                                                                        Gaps
and permit target amplification without PCR, increasing fidelity. The ability to separate desired double stranded targets with allelic selectivity, with or without contemporaneous detection, offers significant advantages over current genotyping methods. The present sequence is an oligonucleotide of the invention.
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Pred. No. 2.9e+02;
0; Mismatches 1; Indels
                                                                                                                                                   2.9%; Score 12.4; DB 1; Length 15; 92.9%; Pred. No. 2.9e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; DNA amplification; ds; mucin-box; G cassette.
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                                                                                                                 Seguence 15 BP; 2 A; 6 C; 5 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU SO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for amplifying a DNA sequence. The current se
box encoding G cassette DNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 5; 33pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mucin-box encoding G cassette DNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JUN-2000; 2000JP-00196242.
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Best Local Similarity 92.9%;
Matches 13; Conservative (
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Best Local Similarity 92.5.
First Local Similarity
First Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified.
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25-MAR-2003 26-JUL-1994

AAQ57378;

RESULT 627 AAQ57378

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The inventors claim an oligo-peptide-carrier conjugate in which the three moieties are covalently linked to one another. The peptide provides a cleavable linker which is cleaved by enzymes which do not degrade antisense oligos (ODNS). The ODN-Largeting ligand linkage must be stable to serum proteases, yet cleaved by the lyosomal enzymes in the target coll. The method involves conjugation of an ODN bearing an electrophilic crosslinking gp. to a peptide which bears two mucleophilic gps of that a nucleophilic handle remains on the peptide. This gp. is used to turther attach the lyosomotropic carrier to the peptide portion of the ODN-peptide conjugate. The peptide is therefore aslo used as a heterobifunctional linker. Two different model ODNs were used - ODNI and CODN-peptide is therefore aslo used as a heterobifunctional linker. Two different model ODNs were used - ODNI and transcript for the Hepatlits B surface antigen in Hep3B cells. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New covalently linked conjugates of oligo:nucleotide, peptide and carrier - utilising surfactant, poly:amine or targetting ligand as lyso somotropic drug carrier.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.9%; Score 12.4; DB 1; Length 16; 92.9%; Pred. No. 3.4e+02; ve 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleotide sequence of the RTBV PCR primer 1.
surface antigen; Hep3B cells; ss.
                                                                                                               /*tag= a
/label= H2N-(CH2)6-O-PO2-
/note= "modified site"
                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Page 19; 77pp; English.
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                                                                                                                                                                                                                                                                           93WO-US012246.
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Les 13, Conservative
                                                                                                                                                                                                                                                                                                                                                       (MICR-) MICROPROBE CORP
                                                                                                                                                                                                                                                                                                                                                                                                 Gall AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1994-217541/26
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                                                                                              misc feature
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                                                                                                                                                                                                                                     23-JUN-1994.
  Hepatitis B
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                                        Synthetic.
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Matches
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                                                                                                                                                                                                                                                            inhibitor; inhibition; ribozyme; treatment; prevention; psoriasis; asthma; inflammatory diseases; cardiovascular condition; hypertension; arthritis; restenosis; angiotensin converting enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic RNA molecules which cleave mRNA - used to treat or prevent inflammatory, arthritic, stenotic or cardiovascular diseases or
                                                                                                                                                                                                                                         specific; cleavage; target RNA; protein; prophylaxis; expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5'-hexylamine modified antisense oligo (ODN1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16 BP; 5 A; 5 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                    Enzymatic RNA molecule ACE mRNA target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide; ODN; modified oligo;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 23; 65pp; English.
                                                               AAQ57378 standard; mRNA; 16 BP.
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92US-00987132.
92US-00989848.
92US-00989849.
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(first entry)
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                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Draper KG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1994-048853/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sullivan SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
02-MAR-1995
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19-JAN-1993;

conditions.

AAQ68223;

RESULT 628

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17-JUL-1992; 07-DEC-1992 07-DEC-1992 17-DEC-1992

Synthetic.

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Gaps

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This is the nucleotide sequence of a PCR primer used in the amplification of the Rice tungro bacilliform virus (RTBV) promoter. The isolated genome-length transcript promoter from RTBV is used for driving gene expression in the vascular bundles of graminaceous plants, especially rice, especially where the gene encodes a protein conferring a desired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotides encoding full-length polypeptides, e.g. secretory and/or membrane proteins, useful for developing medicines for diseases in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; Anti-inflammatory; Osteopathic; Neuroprotective; Nootropic; Gene Therapy; human; secretory protein; membrane proteins; cancer; inflammatory disease; PCR; primer;
                                                                                                                                                                        Rice tungro bacilliform virus promoter - for driving gene expression in vascular bundles of plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ishii S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Otsuki T, Wakamatsu A, Sato H, Ish
Hio Y, Otsuka K, Nagai K, Irie R,
Otsuka M, Nagahari K, Masuho Y;
                                                                                                                                                                                                                                                                                                                                                             2.9%; Score 12.4; DB 1; Length 16; 92.9%; Pred. No. 3.4e+02; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                      Sequence 16 BP; 7 A; 3 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human protein-related PCR primer, SEQ ID 3325.
                                                                                                                                                                                                                 Disclosure; Col 3; 12pp; English
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RES ASSOC BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADASS757 standard; DNA; 16.BP.
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24-JAN-2002; 2002US-0350435P.
                                      91US-00789738.
                                                                91US-00789738
                                                                                                                      Bhattacharyya M;
                                                                                                                                                                                                                                                                                                                                                                                                                       397 AGAAGGICTICTAC 410
                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AGAAGATCTTCTAC 14
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                                                                                                                                                                                                                                                                                                                                                                                             13; Conservative
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J, Isono Y, H
Yoshikawa T, O
                                                                                          UNIW ) UNIV WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-395539/38.
                                                                                                                                              WPI; 1998-582649/49.
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Best Local Similarity
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                                                                  08-NOV-1991;
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Yamamoto J,
            20-OCT-1998,
                                                                                                                      Beachy RN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA55757;
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(REAS-)
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Gaps

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Tamechika I;

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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A partent (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (ft-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (ftk-1) (e.g. tumour angiogenesis, coular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX7575 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; trumour anglogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor; foetal liver kinase 1; ss.
                                                                                                                                               present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecule modulating VEGF receptor(s) gene expression or stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                             Gaps
which the gene is involved, or as target molecules for gene therapy
                                                                    The present invention relates to novel human secretory or membrane proteins (ADA54072-ADA55710) and their coding sequences (ADA52433-ADA54071). The coding sequences are useful in the gene therapy of diseases caused by abnormalies of the proteins, e.g. cancer, inflammatory diseases, osteoporosis or neurological disease. The prequence was used in an example from the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse flt-1 VEGF receptor hammerhead ribozyme substrate #647
                                                                                                                                                                                                                                      Query Match
2.9%; Score 12.4; DB 1; Length 16;
Best Local Similarity 92.9%; Pred. No. 3.4e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                      Sequence 16 BP; 2 A; 5 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stinchcomb D,
                                       Example 8; Page 111; 205pp; English.
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AAX75119 standard; RNA; 17 BP.
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96US-00584040.
                                                                                                                                                                                                                                                                                                                  AGAGAACTCGGTGG 225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                    14 ACAGAACTCGGTGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-259017/23.
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11-JAN-1996;
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ö Gaps ö Score 12.4; DB 1; Length 17; Pred. No. 3.8e+02; 0; Mismatches 1; Indels Sequence 17 BP; 1 A; 8 C; 2 G; 0 T; 6 U; 0 Other; 2.9%; Query Match
Best Local Similarity 92.2.
Best Local 3; Conservative g

1 GGCCAGGAGTGAAA 14

15 GGCCAGGAGTGAGA 2

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AAZ24186 standard; DNA; 17 BP. 03-FEB-2000 AAZ24186; RESULT 632 AAZ24186/c

(first entry)

Human BRCA2 primer scorpion B2731 fragment 1.

Detection; genomic DNA variation; inherited disease; microbial infection; hybridisation; primer; ss.

Synthetic

Homo sapiens

3B2338301-A.

15-DEC-1999

98GB-00025698. 5-NOV-1998;

98GB-00012768. 13-JUN-1998;

(ZENE ) ZENECA LID.

Theaker J, Whitcombe DM; Gibson NJ, Little S,

4PI; 2000-016019/02

Detecting nucleic acids for the diagnosis of heritable genetic disorders and for the detection of microbial organisms in food and biological

Example 7; Page 25; 74pp; English.

This invention describes a novel method (I) for detecting nucleic acids using novel primers and an integrated signaling system. (I) may be used for the detection of variations genomic DNA samples (e.g. from humans, animals and plants). It is particularly useful for detecting inherited diseases (by detecting abnormalities in DNA from patients) and microbial infections (e.g. human immundeficiency virus (HIV) and Hepatitis C infections (e.g. human immundeficiency virus (HIV) and Hepatitis C infections (or bacterial infections of food). (I) provides high levels of sequence specificity, detection sensitivity and high rates of signal applicity and allowing enhanced specificity based on the ready availability of a target binding region (TargBR) for hybridization with product is the target binding region (TargBR) for hybridization with product is the target species so the output signal obtained is directly related to the amount of extended primer. (I) is not dependent on additional hybridization events or ensymmatic steps intra- and inter-strand competition for the probe site is limited so the probe design is simplified and probes which fail to bind under standard assay conditions in spearate probe formats may function in (I). Additionally, homogenous assay formats may be derived from (I). Finally, as the interaction is cycling rates. AAZ24184-Z24190 represent primers used in the method of the propertion. 

Sequence 17 BP; 5 A; 1 C; 8 G; 3 T; 0 U; 0 Other;

2.9%; Score 12.4; DB 1; Length 17;

Query Match

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Gapa

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Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels

Sequence 17 BP; 3 A; 8 C; 1 G; 5 T; 0 U; 0 Other;

the invention

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This invention describes a novel method (I) for detecting nucleic acids using novel primers and an integrated signaling system. (I) may be used using novel primers and an integrated signaling system. (I) may be used an impact of variations genomic DNA samples (e.g. from humans, and plants). It is particularly useful for detecting inherited diseases (by detecting abnormalities in DNA from patients) and microbial infections (e.g. human immunodeficiency virus (HIV) and Hepatitis C infections (e.g. human immunodeficiency virus (HIV) and Hepatitis C viruses or bacterial infections of food). (I) provides high levels of sequence specificity, detection sensitivity and high rates of signal applicity and allowing enhanced specificity based on the ready aniphlicity and allowing enhanced specificity based on the ready aniphlicity and allowing region (TargBR) for hybridization with product is the target binding region (TargBR) for hybridization with product is the target species so the output signal obtained is directly related to the amount of extended primer. (I) is not dependent on product is the target species of the output signal obtained is directly related to the amount of extended primer. (I) is not dependent on strand competition for the probe site is limited so the probe design is simplified and probes which fail to bind under standard assay conditions in spearate probe formats may be derived from (I). Finally, as the interaction is unimplecular, the signal reaction is very rapid, permitting increased contents of the propersent primers used in the method of the broadent of the primer.
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                                                                                                                                                                                                                                                                                                                                                   Detection, genomic DNA variation, inherited disease, microbial infection;
hybridisation, primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting nucleic acids for the diagnosis of heritable genetic disorders and for the detection of microbial organisms in food and biological
                     Gaps
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                     Indels
Pred. No. 3.8e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Theaker J, Whitcombe DM;
                                                                                                                                                                                                                                                                                                                    Human BRCA2 quencher primer B4249.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98GB-00012768
    92.98;
                                                                                                                                                                                                  AAZ24188 standard; DNA; 17
                                                           74
                                                                                                                                                                                                                                                                              (first entry)
  Best Local Similarity 92.9
Matches 13; Conservative
                                                             61 AGTCTCTGCACTAC
                                                                                              16 Acrererecaerae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-016019/02.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                        AAZ24188;
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Page 305
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2 ACTCTCTGCACTAC 15
61 AGTCTCTGCACTAC 74
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ABK00290 standard; RNA; 17 BP (first entry) 12-MAR-2002 ABK00290; 

Human NOGO Hammerhead Ribozyme #290.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; D20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; lymphocytic leukaemia; MCL; immunocytoma; IMC; immune thrombocytopaemia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy:induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creuzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens Synthetic. WO200159103-A2.

16-AUG-2001.

09-FEB-2001; 2001WO-US004273.

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

RIBO-) RIBOZYME PHARM INC. BLATT L. MCSWIGGEN J. CHOWRIRA B M. (BLAT/) 1 (MCSW/) 1

CHOM/)

Chowrira BM; Blatt L, Mcswiggen J,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 70; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NXN motif) prossessing an NCH motif), a G-cleaver (cleaving RNA with a NXN motif), a closed to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2+. The cell and treat a parient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma (NHL), bulky low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky low-grade or follicular non-Hodgkin's lymphoma (NHL), lymphocytic leukaemia, HIV (human immunodeficiency virus) associated NHL, mantle-cell

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CC lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immune thrombocytopaenia, and inflammatory arthropathy. The NOGO-computed immune thrombocytopaenia, and inflammatory arthropathy. The NOGO-computed in the Inflammatory arthropathy. The NOGO-computed in the Inflammatory arthropathy. The NOGO-computed in the Inflammatory acadimated NOGO activity of the conclained may be concacted with the level of NOGO. The treatment may further computes the use of one or more therapies. In particular, the NOGO-cargetting nucleic acid may be used to therapies. In particular, the NOGO-cargetting nucleic acid may be used to therapies. In narvous system (CNS) injury and carebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), chemotherapy-induced neuropathy, and/or other neurodegenerative disease cuasular dystrophy, and/or other neurodegenerative disease cuasular dystrophy, and/or other neurodegenerative disease construction is a hammerhead ribozyme of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; ID20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; lymphocytic leukaemia; MCJ; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arbiropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; cerebroteapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia, Huntington's disease; ceutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                ô
                                                                                                                                                                                                                                                                                                                                                                                 Score 12.4; DB 1; Length 17;
Pred. No. 3.8e+02;
0; Mismatches 1; Indels (
                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 3 A; 6 C; 1 G; 0 T; 7 U; 0 Other;
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28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                                                                                                                                                                                                                                                                                        2.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              288 AAGCTGGTGAAGGA 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 AAACTGGTGAAGGA 4
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Best Local Similarity 92.9%
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J.
(CHOW/) CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-607195/69.
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Synthetic.
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ID ABK(
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constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 132; 200pp; English.

19-FEB-2001; 2001WO-US004273

WO200159103-A2. 16-AUG-2001

Synthetic.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motili), a G-cleaver (cleaving RNA with an NGM with a NGM with an NGM with an NGM with an NGM with an NGM with a reach an amberzyme (cleaving RNA with an NGM with a call to reduce CD20 acidity of the cell and treat a patient having a condition associated with the level of CD20 in the presence of a divalent cation that is preferably NG<sup>2</sup>+.

Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to cleave RNA associated NH, mantle-cell lymphoma (NLI), immunocytoma (INC), small B-cell lymphocytic lymphocytic lymphoma (NLI), immunocytoma (INC), small B-cell lymphocytic lymphocytic contacted with a cell to reduce NOGO gene in the invented and treat a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the NOGO. The treatment may further comprise the use of one or more contender a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the COGO activity of the NOGO. The treatment may further comprise the use of one or more companies. All and treat a patient having a condition associated with the level of therapies. In particular, the NOGO-targetting nucleic acid may be contacted with a cell to reduce NOGO activity of the COGO activity of the

Sequence 17 BP; 1 A; 9 C; 6 G; 0 T; 1 U; 0 Other;

1; Indels 0; Gaps 2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; Live 0; Mismatches 1; Indels Query Match
Best Local Similarity 92.5"
Est Local 3 Conservative

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ABK01168 standard; RNA; 17 BP. ABK01168; RESULT 636 ABK01168/ 

12-MAR-2002 (first entry)

Human NOGO Inozyme #438.

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; D20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lumphoma; leukaemia; human immunodeficiency virus; HIV associated NHL; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHL; martle-cell lymphoma; MCI; immunocytoma; IMC; immune thrombocytopaemia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Homo sapiens

RESULT 637 ABK02396

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGO2). The regulates expression of a neurite growth inhibitor gene (NGO2). The collain and in nozyme (an endolytic nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a contacted may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a nucleic acids (e.g. a ribozyme or a nucleic acids and preserved the noil of the preserved of CD20-targetting nucleic acid is used to cleave RNA with a YGY mortif). The CD20-targetting nucleic acid is used to cleave RNA of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more contacted with the CD20 targetting nucleic acid may be used to the call and treat (NRL), bulky low-grade or follicular non-thodakin's lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-thodakin's lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-thodakin's lymphoma (MRL), immunodeficiency virus) associated MHL mantle-cell lymphoma (MCD), immunodeficiency virus) associated MHL mantle-cell cargetting nucleic acid may be contacted with a cell to reduce NOGO gene in the cargetting nucleic acid may be contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell to reduce NOGO gene in the contacted with a cell and treat a patient having a condition associated with the level of the contacted with a cell and divalent adisable. All pat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence is an inozyme of the invention
                                                                                                                                                                                                                                                                                                                                                                                                          Chowrira BM;
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                                                                                                                                                                                    11-FEB-2000; 2000US-0181797P.
28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
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                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC.
(BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J.
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16 AAACTGGTGAAGGA 3
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Wed Apr 21 12:58:21 2004

ABK02396 standard; RNA; 17 BP.

ABK02396;

12-MAR-2002 (first entry)

Human NOGO Amberzyme #68.

cerebroprotective; nootropic, neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NGO; hammerhead ribozyme; DNAzyme; inozyme; G-Cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; MCL; immunodeficiency virus; HIV associated NHL; mantie-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cenebroty accident; CMA; Alzheimer's disease; multiple sclerosis; chemocherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; Creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease. Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic;

Homo sapiens

Synthetic

WO200159103-A2.

16-AUG-2001

11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J. (CHOW/) CHOWRIRA B M.

Chowrira BM; Blatt L, Mcswiggen J,

WPI; 2001-607195/69.

Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and central nervous system injury.

Claim 88; Page 131; 200pp; English.

The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down cregulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acid cleaving a an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule CG possessing an NCH mocif,), a C-leaver (cleaving RNA with a NGN withen) RNA with a NGN triplet), a zinzyme (cleaving RNA with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD2 or treatment may further comprise the use of one or more cherapies. In particular, the CD20 targetting nucleic acid may be used to cloilicular NHI, lymphocytic cleaksemia, HIV (human immunodeficiency virus) associated NHI, mantle-cell lymphoma (MCL), immunocytoma (IMC), small B-cell lymphocytic lymphoma, confine a second and the NGO gene in the creatment may be contacted with a cell to reduce CD3 gene in the collacted a patient having a condition associated with the level of the NGO. The treatment may further comprise the use of one or more

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therapies. In particular, the NOGO-targetting nucleic acid may be used treat central nervous system (GNS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy induced neuropathy, amyotrophic lateral sclerosis (ALS), Parkinson's disease, ataxia, Huntingcon's disease, Creutzfeldt-Jakob disease, anscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1012.
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                                                                                                                                                                Score 12.4; DB 1; Length 17;
Pred. No. 3.8e+02;
0; Mismatches 1; Indels n
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                                                                                                               sequence is an amberzyme molecule of the invention
                                                                                                                                              Sequence 17 BP; 1 A; 9 C; 6 G; 0 T; 1 U; 0 Other;
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27-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0234587P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US0006663.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
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                                                                                                                                                                              2.9%;
Local Similarity 92.9%;
Nes 13; Conservative (
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                                                                                                                                                                                                                                             307 GCCCCGGGGACCGC 320
                                                                                                                                                                                                                                                                           2 GCCCCGGGGACCCC 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200192524-A2.
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                                                                                                                                                                                                                                                                                                                                                                                            ABN01020;
                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                              RESULT 638
                                                                                                                                                                                                                Matches
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WPI; 2002-179446/23.

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nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for carperssing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as character probes for surface-enhanced laser desorption ionisation, as production, and in vacatines or for replacement therapy. The production, and in vacatines or for replacement therapy. The production associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. HGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO cat fig. wipo.int/pub/published_pot_sequence
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Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
Sequence 17 BP; 8 A; 1 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                           204 GTGAAAGCAGAAA 217
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0; Gaps

Human GDMLP-1 17-mer scanning SEQ ID NO:4 sequence SEQ ID NO:1011. ABN01019 standard; DNA; 17 BP. (first entry) 29-MAY-2002 ABN01019; RESULT 639 ABN01019 

Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.

Homo sapiens.

WO200192524-A2.

06-DEC-2001.

25-MAY-2001; 2001WO-US016981

26-MAY-2000; 2000US-0207456P.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000US-0236359P.
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 201WO-US000663.
30-JAN-2001; 201WO-US000664.
30-JAN-2001; 201WO-US000665.
30-JAN-2001; 201WO-US000666.

30-JAN-2001; 2001WO-US000668 30-JAN-2001; 2001WO-US000669 30-JAN-2001; 2001WO-US000670 05-FEB-2001; 2001US-0266860P

Shannon ME; Chen W, DR, Rank DK, Hanzel Penn SG, Ji Y, Gu Y,

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The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 concleic acids can be used as probes to detect, characterise and quantify mucleic acids can be used as probes to detect, characterise and quantify concleic acids can be used as probes to detect, characterise and quantify convide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therefore the patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration and skeletal muscle disorders. hGDMLP-1 may be used for diagnosing a closyntherapy should skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence data for this patent did not form part to the printed concentration, but was obtained in electronic format directly from MIPO concentration are the present invention.
                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 8 A; 1 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                        Disclosure, SEQ ID NO 1011; 214pp; English.
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Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels Human POSHL1 scanning oligonucleotide SEQ ID NO 1823. ABV91110 standard; DNA; 17 BP. 204 GTGAAAGCAGAA 217 2 GGGAAAGCAGAGAA 15 (first entry) 23-DEC-2002 ABV91110; à ď

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Gaps

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Human, POSHL 1; SH3 domain, POSH-like signalling protein 1, oncoge Rho GTPase, signal transduction, gene expression, cancer, vaccine, gene therapy, transgenic, ss. 30-JAN-2001; 2001WO-US000663. 30-JAN-2001; 2001WO-US000664. 30-JAN-2001; 2001WO-US000665. 30-JAN-2001; 2001WO-US000666. 30-JAN-2001; 2001WO-US000667. 30-JAN-2001; 2001WO-US000669. 30-JAN-2001; 2001WO-US000669. 28-JAN-2002; 2002EP-00001165 Homo sapiens EP1239051-A2 11-SEP-2002. 

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABB8399), a sequence having 65% sequence identity to (SI), CC (SI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-encogene/oncogene product that functions as an adaptor protein that interacts with Rh0 family small GTPsses as well as communitying a specific binding partner. (I) and mucleic acids (II) for identifying a specific binding partner. (I) and mucleic acids (II) concepting (I) are useful for diagnosing, monitoring disease and treating caused by altered axpression of human POSHL1 including diagnosing and cruseful in gene therapy. (II) is useful for conservating microarrays which are useful for measuring and for surveying gene expression and creating cransgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomout of torm part of the present sequence of the present sequence did not form part of the priment by the European Patent Office
                                                                                                                                                                                                                                                                            Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL
-1, useful for treating disorders associated with decreased expression or
activity of human POSHL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human, POSHL 1, SH3 domain, POSH-like signalling protein 1; oncogene; Rho GTPase; signal transduction; gene expression; cancer; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 2.9%; Score 12.4; DB 1; Length 17; Local Similarity 92.9%; Pred. No. 3.8e+02; es 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                     Example 2; SEQ ID NO 1824; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 17 BP; 4 A; 7 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV91108 standard; DNA; 17 BP.
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
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                   30-JAN-2001; 2001WO-US000670.
23-MAY-2001; 2001US-00864761.
10-OCT-2001; 2001US-0328205P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14 GAGGGTCTCTGCA 1
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                                                                                                                            (AEOM-) AEOMICA INC
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                                                                                                                                                                                  Shannon M;
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Best Local S
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ABV91108/c
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                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHI 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3999), a sequence having 65% sequence of 6711, (SI) having 95% deviations, especially conservative substitutions or a cids (SI) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHI 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and mucleic acids (II) for identifying a specific binding partner. (I) and mucleic acids (II) caused by altered expression of human POSHI including disease and treating caused by altered expression of human POSHI including disposing and treating caused by altered expression of numerous constructing microarraps which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to be present by the European Patent Office
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                                                                                                                                                                                                           Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                    Example 2; SEQ ID NO 1823; 60pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000666.
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     10-OCT-2001; 2001US-0328205P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          57 GAGGAGTCTCTGCA 70
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                                                                                                                                                              WPI; 2002-684061/74.
                                                        (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP1239051-A2.
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                                                                                                              Shannon M;
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RESULT 641

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Gaps ô

Shannon M;

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABBB399), a sequence having 65% sequence identity to (S1), (C) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-cncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful of identifying a specific binding partner. (I) and nucleic acids (II) contidentifying a specific binding partner. (I) and nucleic acids (II) consecut in general protein of the signal transduction pathway. (II) is useful for diagnoshing, monitoring disease and treating caused by altered expression of human POSHL1 including diagnoshing and cuseful in the development of vaccines and (II) is useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples continue specification, but is based on sequence information supplied to between by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.88+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; SEQ ID NO 1822; 60pp + Sequence Listing; English.
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              30-JAN-2001, 2001MO-US000664.
30-JAN-2001, 2001MO-US000665.
30-JAN-2001, 2001MO-US000666.
30-JAN-2001, 2001MO-US000666.
30-JAN-2001, 2001MO-US000669.
30-JAN-2001, 2001MO-US000670.
23-MAY-2001, 2001US-00864761.
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ID ABL31374 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-684061/74.
                                                                                                                                                                                                                                                                                                    (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                         Shannon M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino acids (SI, ABBB3999), a sequence having 65% sequence of 6711, (S1) having 95% deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) caused by altered expression of human POSHL1 including disease and treating caused by altered expression of human POSHL1 including microarraps which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The creating cancel is sequence is that of a scanning oligonucleotide useful in examples of present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to berwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                    Novel human SH3 domain (POSH)-like signaling protein 1 polypeptide, POSHL-1, useful for treating disorders associated with decreased expression or activity of human POSHL1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
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2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; SEQ ID NO 1821; 60pp + Sequence Listing; English.
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30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US0006770.
33-MAX-2001; 2001WG-08006770.
10-OCT-2001; 2001US-0328205P.
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06-DEC-2001

28-JAN-2002; 2002EP-00001165.

11-SEP-2002

sapiens 3P1239051-A2

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23-DEC-2002

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ABT39199,
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                                                                                                                                                                                      Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                     Ichihara T, Matsumura Y, Moriya S, Nishida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis; cellular degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 3 A; 3 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human tumour suppressor sequence #2172.
                                                                                                                                                                                                                                                         Claim 10; Page 257; 345pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amson R;
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01-JUN-2001; 2001WO-JP004662.
                                   01-JUN-2000; 2000JP-00164798.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60 GAGTCTCTGCACTA 73
                                                                 (NISN ) NISSHINBO IND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Best Local Similarity 92.9°
Matches 13, Conservative
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                                                                                   SYST-) SYSTEM RES INC
                                                                                                                     Inoko H, Kagiya T,
                                                                                                                                                       WPI; 2002-122074/16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-DEC-2002.
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The invention relates to a novel isolated 17 mer nucleic acid sequence, option in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that they indicate to them under highly stringent conditions, or the complement of any of them, or the complement conditions, or the complement acids of the invention are useful as probes and primers for detecting, indentifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors concaining the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic, virucide, neuroprotective, nootropic, neuroleptic, gene chip, antisense, sense, tumour, cell degeneration, cancer, Alzheimer's disease, schizophrenia, protein chip, gene therapy, tumour suppression, human thkutin, ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                       This sequence represents an isolated mucleic acid sequence associated with tumour suppression or regersation, apoptocasis or virus resistence. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to disapnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
disease, development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 U; 0 Other;
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                                                                                        Claim 1; Page 542; 798pp; French.
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diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polymucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; dc_claaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; coloctal cancer; brain cancer; osophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cosphageal cancer; stomach cancer; bladder cancer; pancreatic cancer; ocervical cancer; stomach cancer; overian acancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chichosphamide; doxcubin; fluorouracall carboplatin; datrexate; cyclophosphamide; doxcubin; fluorouracall carboplatin; datrexate; reclenated arthritis; restences; crohn's disease; asthma; diabetes; rheumatoid arthritis; restences; lunguals; multiple sclerosis; schaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; seppais; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
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94US-00245466.
94US-00291932.
96US-00777916.
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MCSW/) MCSWIGGEN J.
DRAP/) DRAPER K G.
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23-DEC-1996;
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regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NERD), where (I) is an inozyme, Zinzwe, G-Cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treathing a parient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of advalent cation, especially MG·2+. The enzymatic and antiense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, ossophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as the control method of the color o
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85.7%; Pred. No. 3.8e+02;
ive 1; Mismatches 1; Indels
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nes 12; Conservative
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18-MAY-1994;
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23-DEC-1996;
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(STIN/) STINCHCOMB D T.

23-MAY-2001; 2001US-00864785

US2002177568-A1 Homo sapiens

28-NOV-2002

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Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                   Stinchcomb DT, Mcswiggen J, Draper KG;
                                                                          Claim 3; Page 50; 72pp; English.
MCSW/) MCSWIGGEN J.
                                 WPI; 2003-340953/32.
      (DRAP/) DRAPER K G.
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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzwe, g-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or milidicug resistant cancer. The method involves use of other drug cervical, esistant cancer. The method involves use of other drug chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, canid molecules are also useful for treating inflammatory disease such as acid molecules are also useful for treating inflammatory disease such as colection, gene therapy applications, ischemials, traparison injury celectrian, mis accordance in the properties of a novel paramatic constants. infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule

Seguence 17 BP; 3 A; 4 C; 8 G; 0 T; 2 U; 0 Other;

0; Gaps 2.9%; Score 12.4; DB 1; Length 17; 85.7%; Pred. No. 3.8e+02; ive 1; Mismatches 1; Indels 1; Indels 12; Conservative Similarity Query Match Local

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ACA06441 standard; RNA; 17 BP 03-JUN-2003 ACA06441; RESULT 649 ACA06441/c 

(first entry)

NFKB sub-unit modulating inozyme substrate #260.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; coesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; hadder cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubn; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn; G disease; besity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/garaft rejection; reperfusion injury; glomerulonephritis; allergic alrway inflammation; inflammatory bowel disease; infection; ss.

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regulates expression of a sequence encoding a subunit of muclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat carder and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for clearing RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, creating resistant cancer. The method involves use of other drug cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug cervical, head and neck, ovarian tanibodies REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, conditable or radiation flerapy. The enzymatic and antisense nucleic conditable or radiation therapy. The enzymatic and antisense nucleic conditable or radiation therapy. The enzymatic and antisense nucleic conditable or radiation therapy. The enzymatic and antisense nucleic conditable or radiation therapy. The enzymatic and antisense nucleic conditable or adiation therapy. The enzymatic and antisense such as reals useful for treating inflammatory disease such rejection, gene therapy applications ischemial/trepartusion injury inflammatics, restenosis, intended to central nervous system (CNS) and myocardial), glomerulonephritis, inflammatic, separation, miliammatory for the content of th
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94US-00291932.
96US-00777916.
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Best Local Similarity 92.9
Matches 13; Conservative
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(MCSW/) MCSWIGGEN J.
(DRAP/) DRAPER K G.
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15-AUG-1994;
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33-DEC-1996; 

G-cleaver, amberzyme, cancer; REL-A activity, breast cancer; human; lung cancer; prostate cancer; REL-A activity, breast cancer; human; lung cancer; prostate cancer; cancer; brain cancer; human; cesophageal cancer; stomach cancer; braid cancer; paroreatic cancer; cervical cancer; and and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemocherapy; paclitaxel; docetaxel; cipplatin; methotrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; isofhamia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss. Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785.

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 07-DEC-1992; .8-MAY-1994;

STIN/) STINCHCOMB D T.

DRAP/) DRAPER K G. MCSWIGGEN MCSM/)

Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 55; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down cregulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFKB), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for deaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-t. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, corrical, head and neck, ovarian cancer, melanoma, lymphoma, glicma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal attibodies; REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, concentral nervous system (CMS) and mycardial), glomeruloned second arthritis, restenosis, asthma, Crohn's disease such as crejection, gene therappy applications, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic contral nervous system (CMS) and mycardial), glomeruloned is a novel enzymatic nucleic acid molecule

Sequence 17 BP; 0 A; 9 C; 7 G; 0 T; 1 U; 0 Other;

Gaps . 0 2.9%; Score 12.4; DB 1; Length 17; ilarity 92.9%; Pred. No. 3.88+02; Conservative 0; Mismatches 1; Indels Query Match Best Local Similarity Matches 13; Conserva

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ACA06442 standard; RNA; 17 RESULT 651 ACA06442,

BP.

ACA06442;

(first entry) 03-JUN-2003

NFKB sub-unit modulating inozyme substrate #261.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; prain cancer; colorectal cancer; pancreatic cancer; cervical cancer; pancreatic cancer; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotraxate; cyclophosphamide; docetaxel; cisplatin; methotraxate; cyclophosphamide; docetaxel; cisplatin; dispecific inhibitor; cyclophosphamide; docetaxel; cisplatin; displatin; diabetes; gemoitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.

Homo sapiens.

US2002177568-A1.

28-NOV-2002.

23-MAY-2001; 2001US-00864785

92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916. 07-DEC-1992;

18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

STIN/) STINCHCOMB D T. MCSW/) MCSWIGGEN J. (DRAP/) DRAPER K G.

Draper KG; Stinchcomb DT, Mcswiggen J,

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 31; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor cappa B NPKED, where (I) is an inozyme, 2inzyme, 6-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (C) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or miltidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphande, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphande, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphande, doxorubin, fluorouracil carboplatin, adatesate, cole gemoitable or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as theumatoid arthritis, restencesis, asthma, Crohn's disease, diabetes,

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obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graf rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic
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                                                                                                                                                                        Query Match
2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                         Sequence 17 BP; 1 A; 11 C; 3 G; 0 T; 2 U; 0 Other;
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94US-00245466.
94US-00291932.
96US-00777916.
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                                                                                                 nucleic acid molecule
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(DRAP/) DRAPER K G.
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15-AUG-1994;
23-DEC-1996;
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cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.

(I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MS-2+. The enzymatic and antisense mucleic acid molecules are useful for traating breast, lung, prostate, colorectal, brain, oseophageal, stoomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or themotherapy including paclitaxel, docetaxel, cisplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide doxorubin, fluorouracil carboplatin, edatrexate, cand molecules are also useful for treating inflammatory disease such a caid molecules are also useful for treating inflammatory disease, cobesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft refection, gene therapy applications, ischaemia/reperfusion injury celection, gene therapy applications, ischaemia/reperfusion injury central nervous system (CNS) and myocardial), glomerulonephritis, infinection. This sequence represents the substrate of a novel enzymatic connection desiral mervous system (SNS) and myocardial).
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Pred. No. 3.8e+02;
1; Mismatches 1; Indels
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94US-00245466.
94US-00291932.
96US-00777916.
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Best Local Similarity 85.7%;
Matches 12; Conservative 1
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MCSWIGGEN J.
DRAPER K G.
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18-MAY-1994;
15-AUG-1994;
23-DEC-1996;
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(DRAP/)
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Shannon M, Gu Y, Nguyen C;

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The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor regulates expression of a sequence encoding a subunit of nuclear factor cancer and is useful for down-regulating REI-A activity in a cell, for cancer and is useful for down-regulating REI-A activity in a cell, for creating a pattent having a condition associated with the level of REI-A. (I) is useful for cleaving RNA comprising a sequence of REI-A gene, in the presence of a divalent cation, especially Mg^2+ The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or cartical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or ultidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REI-A-specific inhibitors or chemotherapy including pacificaxi, fluorouracil catopolatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil catopolatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil catopolatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil catopolatin, and antisense nucleic acid molecules are also useful for treating inflammatory disease such as repeated conformance disease, lupus, multiple sclerosis, transplant/graft cajection, gene therapy applications, ischaemia/reperfusion injury cepterial nervous system (CMS) and myocardial, glomerullonsphritis, sepsis, allergic airway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic inflection. This sequence represents the substrate of a novel enzymatic
                                                                                 Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
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zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
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Mcswiggen J, Draper KG;
                                                                                                                                                                             Claim 3; Page 55; 72pp; English
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Query Match
Best Local Similarity 92.99
Matches 13, Conservative
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                                             WPI; 2003-340953/32.
  Stinchcomb DT,
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ADB00481/c
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ3 is proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 1602.3 and MDZ12, MDZ7, MDZ7, MDZ4, MDZ7, cor in manufacturing a medicament for treating or preventing a disorder or sescoiated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are protein are useful at therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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zinc finger protein; MD21; MD24; MD212; MD212; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
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                                                                                                                         New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                         Example 8; SEQ ID NO 1467; 103pp; English.
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Query Match
Best Local Similarity 92.99
Matches 13, Conservative
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                                                                  WPI; 2003-423107/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB00483;
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ADB00483/c
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0; Gaps

2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; ive 0; Mismatches 1; Indels

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proteins and their coding sequences: MD24, MD24, MD212. MD23 is encoded at chromosome 7q22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 1g21.1, and MD212 is encoded at chromosome 15q26.1. The MD23, MD24, MD27, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder sascoiated with decreased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease acids and proteins are also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapoutle agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                      The present invention relates to novel human zinc finger-containing
Example 8; SEQ ID NO 1469; 103pp; English.
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Sequence 17 BP; 3 A; 6 C; 6 G; 2 T; 0 U; 0 Other;

Gaps ; 0 Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

262 CGGTGCACCTGGAG 275 CGGTGCACCTGCAG 15 ઠે 셤

ADA99414; RESULT 656 ADA99414

Human MDZ3 scanning oligonucleotide SEQ ID 403. BP. ADA99414 standard; DNA; 17 (first entry) 20-NOV-2003 

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ1; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15g26.1; cancer; developmental disorder; ss

Homo sapiens

EP1281758-A2.

JS-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181.

(AEOM-) AEOMICA INC.

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 403; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7922.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15p11.2 and MD212 is encoded at chromosome 15p26.1. The MD23, MD24, MD27, and MD212 sequences are useful in therapy,

or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for disgnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as theraputic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention. 888888888888888

Sequence 17 BP; 2 A; 9 C; 1 G; 5 T; 0 U; 0 Other;

ö Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

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ADA99489 standard; DNA; 17 BP.

ADA99489;

(first entry) 20-NOV-2003 Human MDZ3 scanning oligonucleotide SEQ ID 478.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens.

EP1281758-A2.

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181. 

(AEOM-) AEOMICA INC

Shannon M, Gu Y, Nguyen

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WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 478; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7922.1, MD24 is encoded at chromosome for an encoded at chromosome for in manufacturing at chromosome 15926.1. The MD27 is encoded at chromosome 15926.1. The MD24, MD27, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder or in manufacturing a medicament for treating or preventing a disorder ansociated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The

293 GGTGAAGGACCTGA 306

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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11 2 and MDZ12 is encoded at chromosome 16p12 and MDZ12 is encoded at chromosome 16p2 in manufacturing a medicament for treating or preventing a disorder or associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acides and proteins are also useful for disquesting remonitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ1, or MDZ12. The nucleic acide can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human;
zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1;
chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ10 or MDZ12, e.g. cancer.
                                                                                                                                              Gaps
proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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1 Similarity 92.9%; Pred. No. 3.8e+02;
13; Conservative 0; Mismatches 1; Indels
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                                                               Sequence 17 BP; 3 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Human MDZ3 scanning oligonucleotide SEQ ID 482.
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                                                                                                                                                                                                                                                                                                                            ADA99493 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shannon M, Gu Y, Nguyen C;
                                                                                                                                                                                    292 TGGTGAAGGACCTG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              developmental disorder; ss
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                                                                                                                           Local Similarity
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                                                                                                          Query Match
                                                                                                                                                                                                                                                                                     RESULT 658
                                                                                                                                              Matches
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 7g21.3 MDZ4, MDZ12 and MDZ12 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, 15g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder. Co associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosling or monitoring a disease acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12, energia gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes for detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12, energia cuseful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                    Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                 Human MDZ3 scanning oligonucleotide SEQ ID 1468.
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                                                                                                            BP
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                                                                                                            ADB00482 standard; DNA; 17
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Best Local Similarity 92.9%
Matches 13, Conservative
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GGTGGAGGACCTGA 14
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ADB00484/c
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0; Gaps

Query Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels

ADB00484;

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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-HER2, K-Ras, H-Ras, and HIV acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, shown in NaZS9889 - ABZ62216, ABZ64944 - ABZ65531, ABZ65520 - ABZ65524, ABZ65531 - ABZ65531 - ABZ65524 + Duman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 144; 185pp; English.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                             29-MAY-2002; 2002WO-US016840.
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                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                            WO200297114-A2
                                                                                                           Homo sapiens.
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                                                                                                                                                                                                            05-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ4, MDZ4, MDZ12, MDZ12 mDZ3 is cancoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.2, MDZ7 is encoded at chromosome 6721.2, and MDZ12 is encoded at chromosome 1671.2 and MDZ12 is encoded at chromosome 1671.2 and MDZ12 sequences are useful in therapy, or in manifacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                          Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ1; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23,
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                                                                                                                                                            Human MDZ3 scanning oligonucleotide SEQ ID 1470.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MDZ4, MDZ7 or MDZ12, e.g. cancer.
             ADB00484 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 262 CGGTGCACCTGGAG 275
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Best Local Similarity 92.9°
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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ABZ65139

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ABZ65139

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Gaps

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RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; 88.
Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV;
                                                                                                                                                                                                                       26-MAR-2001, 2001US-00817879.
08-UUN-2001, 2001US-00867478.
08-UUN-2001, 2001US-0256876P.
24-OCT-2001, 2001US-0335059P.
05-DEC-2001, 2001US-0337055P.
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                                                                                                                     Hepatitis C virus
                                                                                                                                               WO200281494-A1.
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                                                                                                                                                                                                                                                                                                                                                                      (PAVC/) (
(LEEP/) I
(DRAP/) I
(ROBE/) I
                                                                                                                                                                                                                                                                                                      (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
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  The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV genes and/or HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and insense states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
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                                                                                                                                                                                                                                                                                                                                                          Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                                                                            Pavco P, Lee
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                                                                                                                                                                                                                                                                                            Mcswiggen J, Morrissey D,
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                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 298; 387pp; English.
                                                                                           08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
                                                      26-MAR-2002; 2002WO-US009187
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                                                                                                                                                             RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                          Roberts E;
                                                                                                                                                                                   MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                            Macejak D,
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                                                                                                                                                                                                                             PAVCO P.
LEE P.
                                                                                                                                                                                                                                                   DRAPER K.
    WO200281494-A1.
                                                                               26-MAR-2001;
                             17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                       infection.
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                                                                                                                                                                                                                                                     (DRAP/)
(ROBE/)
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(PAVC/)
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                                                                                                                                                                        (BLAT/)
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(MCSW/)
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Lee P;

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Mcswiggen J, Morrissey D, Pavco

Roberts E;

Macejak D,

ROBERTS E. LEE P. DRAPER K.

RIBOZYME PHARM INC.

BLATT L. MACEJAK D. MCSWIGGEN J. MORRISSEY D.

PAVCO P.

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, or inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, and enzymatic nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV Genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and methods of the invention and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV invention.
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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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2.9%; Score 12.4; DB 1; Length 17;
Best Local Similarity 92.9%; Pred. No. 3.8e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                     Claim 1; Page 260; 387pp; English.
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                                                                                         infection.
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substrate sequence #1477.

BP

RESULT 663

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Sequence 17 BP; 3 A; 8 C; 6 G; 0 T; 0 U; 0 Other;

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HRV) RNA. The nucleic acid molecules include antisense and erzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, Inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, DNAzymes, are nucleic acid decoy molecules and aptemers that bind to HBV reverse transcriptuse primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV Genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression of HBV methods of the invention and/or replication of HCV. The compounds and methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV increase.
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                                                                                                                                                                                                                                                                                        Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
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                                                                                                                                                                                                                                       HCV minus strand DNAzyme substrate sequence #801
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                                                    ACD62938 standard; RNA; 17 BP
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08-UUN-2001; 2001US-0087478.
24-OCT-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335059P.
05-DEC-2001; 2001US-0337055P.
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Roberts E;
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MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus.
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DRAPER K.
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                                                                                                                     ACD62938;
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RESULT 664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                    Cytostatic; virucide; neuroprotective; noctropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                     Murine oligonucleotide associated with tumour supression, SEQ ID 5492.
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Query March 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                           ACC68245 standard; DNA; 17 BP.
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                                                                            331 CGGACGACCAGGGC 344
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                                                                                                                 3 cceaceaccadesc 16
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                                                                                                                                                                                                                                                                                                                                                                                                                 schizophrenia; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus.
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                                                                                                                                                                                                                                              ACC68245;
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ACC65338
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ACC65338;

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Disclosure; Page 77; 738pp; French.
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| ADB43561 standard; DNA; 17 BP.
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(first entry)
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                   WO2003025176-A2
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04-DEC-2003
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                                                         27-MAR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                        Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.
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                                                                                                                  Murine oligonucleotide associated with tumour supression, SEQ ID 2585
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  ACC65338 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 recreatedaceae 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-333167/31.
                                                                                                                                                                                                                                                                                                WO2003025176-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-2003
                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    relerman A,
                                                                              01-JUL-2003
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27-MAR-2003

Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ss.

ACC63151;

RESULT 667 ACC63151/

Query Match Best Loca Matches

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Mus musculus

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The present invention relates to murine oligonucleotides (ACC62754-ACC68806), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, gene chip) in vitro as antifying nucleic acid, e.g. as one component of a gene chip; in vitro as (anti)sense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or call degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic, antiviral, neuroprotective, nootropic, neuroleptic, ss;
primer, probe, tumour suppression, tumour reversion, apoptosis,
virus resistance, transgenic animals, Alzheimer's disease; schizophrenia;
New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                     specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumour suppression/reversion associated nucleotide #3884.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match 2.9%; Score 12.4; DB 1; Length 17; Best Local Similarity 92.9%; Pred. No. 3.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 3 A; 3 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tuijnder M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-SEP-2001; 2001FR-00011981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-SEP-2002; 2002WO-IB004219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         relerman A,
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WPI; 2003-441574/41. 

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fragments of at least 15 consecutive nucleotides of these nucleotides, a fragments of at least 15 consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisenes sequences, of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing the mand cells containing the vectors), the encoded polypeptides and antibodies of viral infections or diseases characterized by development of tumours or cell degeneration (e.g. Alzhelmer's disease or schizophrenia).

Analysis of the expression of the nucleotides can be used for disapnosis and/or prognosis of these diseases. The nucleotides and polypeptides can and/or prognosis of these diseases. The nucleotides and polypeptides can also be used to screen for their specific interactive molecules.
New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                          Disclosure; Page 486; 771pp; French
```

Sequence 17 BP; 9 A; 1 C; 5 G; 2 T; 0 U; 0 Other;

expression of the nucleotides

Gaps ; 0 Match 2.9%; Score 12.4; DB 1; Length 17; Local Similarity 92.9%; Pred. No. 3.8e+02; les 13; Conservative 0; Mismatches 1; Indels Query Match Best Loca Matches

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ADB45240 standard; DNA; 17 BP. (first entry) 18-DEC-2003 diagnosis. ADB45240; RESULT 669 ADB452 

Tumour suppression/reversion associated nucleotide #5563.

cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss; primer; probe; tumour suppression; tumour reversion; apoptosis; virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;

Homo sapiens.

WO2003040369-A2.

15-MAY-2003.

17-SEP-2002; 2002WO-IB004219.

17-SEP-2001; 2001FR-00011981.

Tuijnder M; Felerman A, Amson R,

(MOLE-) MOLECULAR ENGINES LAB

WPI; 2003-441574/41.

New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related

polypeptide and antibodies.

Disclosure; Page 682; 771pp; French.

The invention relates to the isolation of 6327 mucleotide sequences, fragments of at least 15 consecutive mucleotides of these mucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridises under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the mucleotides are used as probes or primers for detecting, identifying, quantifying and/or amplifying nucleic acids, as in vitro sense and antisense sequences, of nucleotides involved in tumour sequencesion or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as recombinant polypeptides, and to prepare transgenic animals, as recombinant models. The nucleotides (also vectors containing them and calls containing the vectors), the encoded polypeptides and antiboddes (Ab) against the polypeptide are useful for prevention and/or treatment of tumours or call degeneration (e.g. Alzheimer's disease or schizophrenial). Analysis of the expression of the nucleotides can be used for diagnosis and be used to screen for their appearative molecules, and polypeptides can also be useful for treating diseases associated with abnormal expression of the nucleotides 

Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 U; 0 Other;

Gaps ö 2.9%; Score 12.4; DB 1; Length 17; 92.9%; Pred. No. 3.8e+02; tive 0; Mismatches 1; Indels Query Match 2.9 Best Local Similarity 92.9 Matches 13; Conservative

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ADE13461 standard; DNA; 17 BP. (first entry) 29-JAN-2004 ADE13461; RESULT 670 ADE13461/c

HLA class I allele specific primer #77.

ss; primer; PCR; human; Human Leukocyte Antigen; HLA; genotype.

Homo sapiens.

US2003165884-A1.

04-SEP-2003.

20-DEC-1999; 99US-0172768P. 20-DEC-2000; 2000US-00747391. 25-APR-2002; 2002US-00133779. 

(STEM-) STEMCYTE INC. Tonai R; Chow R,

WPI; 2003-874916/81.

Identifying class I or II Human Leukocyte Antigen genotypes using hybridization and amplification assays.

Claim 7; SEQ ID NO 79; 66pp; English.

The invention relates to a method of identifying a class I or II Human Leukocyre Antigen (HLA) genotype of a subject using hybridisation and amplification assay. The method is used for determining the HLA genotype

671 RESULT 67 AAQ22412/

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Human, PRO; EST; expressed sequence tag; PCR primer; hybridisation;
probe; blood coagulation disorder; cancer; cellular adhesion disorder;
secreted protein; transmembrane protein; ss.
                           Human PRO274 PCR forward primer 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                 9805-0079728P-
9805-0079728P-
9805-0079728P-
9805-0080107P-
9805-00801294P-
9805-0080133P-
9805-0080133P-
9805-0080133P-
9805-0080133P-
9805-0080133P-
9805-0080137P-
9805-0080137P-
9805-0080137P-
9805-008120P-
9805-008180P-
9805-008180P-
9805-008180P-
9805-008180P-
9805-008180P-
9805-008180P-
                                                                                                                                                                                                                                                                                                     98US-00040220.
98US-0078886P.
98US-0078910P.
                                                                                                                                                                                                                                                                                                                                          98US-0078936P.
98US-0078939P.
98US-0079294P.
    07-DEC-1999 (first entry)
                                                                                                          Synthetic.
Homo sapiens.
                                                                                                                                               WO9946281-A2
                                                                                                                                                                                                08-MAR-1999;
                                                                                                                                                                       16-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This oligonuclectide was used in an example of synthesis of 3'- acridine-tailed oligonuclectide from acridine-CPG. Blockage of the 3' terminal phosphodiester bond improves resistance to nucleases in serum-contg. media. The new synthesis method avoids the derivatisation step of prior art methods and the possible loss and difficult separation. See AAQ22411-Q22415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Solid support synthesis of 3'-tailedoligo-nucleotide(s) via linker gp. provides nuclease resistant prods. opt. with intercalation to improve anti-sense bonding to DNA or RNA strand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                              Gaps
of a subject. The present sequence represents a HLA class I allele specific primer.
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                                                                                                                                                                                                                                                                                                                     Acridine-CPG; nuclease resistance; controlled pore glass; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.4; DB 1; Length 18;
Pred. No. 4.3e+02;
0; Mismatches 1; Indels
                                                                     Length 17;
                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 18 BP; 3 A; 8 C; 1 G; 6 T; 0 U; 0 Other;
                                         Sequence 17 BP; 2 A; 6 C; 7 G; 2 T; 0 U; 0 Other;
                                                                    Score 12.4; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tabone JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 12; Page 38; 78pp; English
                                                                                                                                                                                                                                                                                               3'-acridine-tailed oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reed MW, Meyer RB, Petrie CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ33902 standard; DNA; 18 BP.
                                                                                                                                                                                                                  AAQ22412 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       90US-00574348.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                              91WO-US006143
                                                                     Query Match
Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           33 TGGGACGAAGATGG 46
                                                                                                                         373 TCCTGGACCGCGAC 386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 rereaceaagares 5
                                                                                                                                                  14 rccreeaccecec 1
                                                                                                                                                                                                                                                                      15-JUL-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MICR-) MICROPROBE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-096825/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-1990;
10-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                            VO9203464-A
                                                                                                                                                                                                                                                                                                                                                                                                      05-MAR-1992
                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ33902
                                                                                                                                                                                                                                              AAQ22412;
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RESULT 672 AAZ33902/C ID AAZ339 XX AC AAZ339

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The present invention describes secreted and transmembrane polypeptides and their polynucleotides. The nucleotide sequences are useful as sources of probes, primers, for chromosome mapping, and for generation of antisense sequences. They can also be used to create transgenic animals. The proteins can be used to treat a variety of diseases and disorders, depending on their function. Diseases that may be treated include blood coagulation disorders, cancers and cellular adhesion disorders. They may also be used to raise antibodies, AA231891 to AA234318, and AAY41685 to AAX41774 represent polynucleotide and polypeptide sequence given in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New secreted and transmembrane polypeptides and their polynucleotides, useful for treating blood coagulation disorders, cancers and cellular adhesion disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Goddard A, Gurney A, Yuan J, Baker KP, Chen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 2.9%; Score 12.4; DB 1; Length 18; Local Similarity 92.9%; Pred. No. 4.3e+02; les 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 4; Page 183; 530pp; English.
                                       9803-0084441P

9803-0084598P

9803-0084637P

9803-0084643P

9803-0084643P

9803-0084643P

9803-0084643P

9803-0085333P

9803-0085333P

9803-0085333P

9803-0085833P

9803-0085833P

9803-008583P

9803-008583P

9803-008583P

9803-008583P

9803-008583P

9803-008583P

9803-008583P

9803-008583P

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9803-00858

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9803-00858
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                                                                                                                                                                                                                                                                                                                                                                                                       98US-0087106P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0100038P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18 GAÁCTCCGTGGCGG S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999~551358/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                22-MAY-1998;
22-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wood WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ91453;
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                                                                                                                                                                                                                                               15-MAY-1
15-MAY-1
                                                                                                                                                            -MAY - 1
-MAY - 1
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15-MAY-1
                                                                                                                                                                                                                                                                             .5-MAY-
                                                                                                                                                                                                                                                                                                                       8-MAY-
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AAZ91453
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Antisense oligonucleotides, useful for inhibiting human Ship-2 expression and for detecting nucleic acids encoding Ship-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes phosphorothioate antisense cligonucleotides that specifically hybridise with, and inhibit the expression of, nucleic acids encoding human Ship-2 (also called SH2-containing phosphatidylinosited phosphatase-2). Also described is a method of inhibiting the expression of Ship-2 in human cells or tissues in vitro comprising contacting the cells with the phosphorothioate antisense oligonuclectides. The phosphorothioate antisense oligonuclectides. The phosphorothioate antisense oligonuclectides to be used to treat animals (especially humans) suspected of having or being prone to a disease or condition associated with Ship-2 expression. The present sequence represents a phosphorothioate antisense oligonucleotide for human Ship-2, from the
                                                                   Human, Ship-2; antisense oligonucleotide; phosphorothioate; detection; inhibition; SH2-containing phosphatidylinositol phosphatase-2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human biallelic marker upstream amplification primer SEQ ID NO:4482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Puman genome; biallelic marker; high density disequilibrium map;
genomic map; haplotype; phenotype; polymorphic base; genotyping;
haplotyping; hybridisation; identification; characterisation;
amplification; single nucleotide polymorphism; SNP; PCR primer;
diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                  Human Ship-2 phosphorothioate antisense oligonucleotide #30735.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            / Match 2.9%; Score 12.4; DB 1; Length 18; Local Similarity 92.9%; Pred. No. 4.3e+02; Local 13; Conservative 0; Mismatches 1; Indels

    .18
    /*tag= a
    /note= "phosphorothioate linkages"

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                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Col 40; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ70126 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                           99US-00339964.
                                                                                                                                                                                                                                                                                                                         99US-00339964.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   273 GAGCAGGGCGCCAC 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 GAGCAGGCAGCAC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-2001 (first entry)
22-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-181819/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention
                                                                                                                                                             Key
modified_base
                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                         25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                          25-JUN-1999;
                                                                                                                                                                                                                                                    US6025198-A.
                                                                                                                                                                                                                                                                                    15-FEB-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ70126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Invention, which contains a polymorphic base at position 24 of their nucleotide sequences. AAX68579 to AAX7740 represent amplification primers for the biallelic markers of the biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the invention man are one, and in complex association studies and haplotyphag studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical as of the analysis of the disease as well as other treatment.

N.B. The SEQ ID NOS 2852, 2913, 2974, 3055, 3096, 3157, 3227, 3297 and and also are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ65654 to AAZ69578 represent human biallelic markers from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, secreted protein; transmembrane protein; PRO; EST; cytostatic; expressed sequence tag; detection; cancer; PCR primer; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 2 A; 7 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO274 forward PCR primer SEQ ID NO:14.
                                                                                                                                                              Cohen D, Blumenfeld M, Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                     Claim 9; Page 2561; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US005028.
99US-0123957P.
99US-0126773P.
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990S-0131445P.
990S-0141037P.
990S-0141037P.
99US-0141037P.
99US-014508P.
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98US-0082614P.
98US-0109732P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        212 AGAGAACTCGGTGG 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-PEB-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                        map of the human genome.
                                                                                                                                                                                                                               WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention
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                                                                                             (GEST ) GENSET.
21-APR-1998;
23-NOV-1998;
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02-DEC-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 676
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       HERE REPRESENTE SERVICE SERVIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention primers for the biallelic markers. The biallelic markers of the invention continue sequences in complex association studies and haplotyping studies which are useful in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions of the targets for the development of pharmaceutical efficacions tresponses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

When the SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel biallelic markers used to construct a high density disequilibrium map of the human genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human biallelic marker downstream amplification primer SEQ 1D NO:10930.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches 1; Indels 0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 1 A; 7 C; 1 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 8; Page 1186; 2745pp; English.
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                 (GEST ) GENSET.
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   Homo sapiens
                                                                WO9954500-A2
                                                                                                                                                                                                                                                                                                23-NOV-1998;
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Query Match

AAZ76574;

RESULT 675 AAZ76574/c

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The present invention describes a method for distinguishing a human leukocyte antigen (HLA) class I antigen or allele by a combination of polymerase chain reaction (PCR) using a primer pair whereby all HLA-A, -B or -C alleles can be amplified or using reverse hybridisation analysis comprising a DNA probe covalently bonded to microtitre plate wells which are hybridisable specifically with the base sequence of at least one specific HLA-A, -B or -C allele. The method is applicable in gene typing, judging donor-recipient compatibility during organ transplant and correlation analysis for diagnosis of various diseases. The method is simple, rapid and accurate, with possibility of mechanisation and automation, without the problems encountered by using the prior-art techniques. AAAA66943 to AAAA6707 represent oligonucleotide probes and PCR primers for use in the method of the present invention
                                                                                  Simple, rapid and accurate method for distinguishing HLA class I allele type with possibility of mechanization and automation, applicable in judging donor-recipient compatibility during organ transplant and disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genetic clustering by distributing members into optimal numbers of clusters determined by a hierarchical clustering algorithm or by pairedpair analysis of homozygous pairs in clusters got from non-hierarchical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cluster; hierarchical clustering algorithm; population based study; clinical trial; DNA fingerprint; genetic profile analysis; PCR primer; SNP; single nucleotide polymorphism; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 5 A; 5 C; 6 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                             Claim 8; Page 66; 83pp; Japanese.
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07-JUL-2000; 2000US-0216897P.
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  Kaneshige T;
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                                             WPI; 2000-400097/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                   diagnosis.
  Moribe T,
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC78458 to AAC78599 represent polynucleotide and EST (expressed sequence tag) sequences which encode secreted or transmembrane PRO polypeptides. The PRO polynucleotides and polypeptides have cytostatic activity. The polynucleotides and polypeptides can be used for detecting the presence of PRO polypeptides can be used for detecting the presence of PRO polypeptides in samples, for linking bioactive molecules to cells for specific targeting biological activities of cells, using the polypeptides for specific targeting. The polypeptide targeting can be used to kill the target cells, e.g. for the treatment of cancers. The polypeptide pairs provide specific targeting of bioactive molecules to cells. AAC78670 to AAC78897 represent PCR primers and probes used in the isolation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                    Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Baton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi CJ, Gurney AL, Hillan KJ; Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human leukocyte antigen, HLA, class I allele type, probe, PCR primer,
amplification, hybridisation, organ transplant, gene typing, diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                         Novel PRO polypeptides and polynuclectides used in detection methods, trarget bioactive molecules to specific cells, and to modulate cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 4; Page 235; 636pp; English.
02-DEC-1999; 99WO-US028565.
16-DEC-1999; 99WO-US0310095.
30-DEC-1999; 99WO-US031274.
30-DEC-1999; 99WO-US031274.
65-JAN-2000; 200WO-US000277.
06-JAN-2000; 200WO-US000377.
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                            (GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   activities.
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RESULT 677 AAA6701

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The present invention relates to a method of determining polynucleotide expression, which comprises hybridising digested cDNA to a capture probe coupled to a solid particle under stringent conditions, where the capture probe is specific for the target polynucleotide and the particle identifies the capture probe. The method is useful for expression profiling, where the presence and/or the amount of a target condition, disorder, or predisposition associated with a change in expression patterns, in determining the developmental or physiological state of a cell or tissue, for detecting SNPs, which may be used to screen individuals for a genetic predisposition to a disease, condition, or disorder, and in marker assisted selection. The present sequence is a hybridisation tag described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining polynucleotide expression, useful for expressing profiling or detecting single nucleotide polymorphisms, comprises hybridizing digested cDNA, to a capture probe coupled to a solid particle under stringent
                             sample, involving applying a hierarchical clustering algorithm to the sample members, determining the optimal number of clusters based on this and distributing the sample members into clusters using non-hierarchical clustering. The methods are useful in population based studies such as clinical trials, DNA fingerprinting and genetic profile analyses. The present sequence was used to demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid analysis, microarray, single nucleotide polymorphism, SNP, multiplex; expression analysis; hybridisation tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Drosophila ubx gene SNP analysis universal hybridisation tag #31.
                   present invention describes methods of clustering members of
                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                       Length 18;
                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                 Seguence 18 BP; 3 A; 4 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                     Score 12.4; DB 1;
Pred. No. 4.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hinkel CA, Kimmerly WJ, Yang L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 34; Page 29; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-2001; 2001US-0264972P.
02-FEB-2001; 2001US-0266186P.
04-JUN-2001; 2001US-0295986P.
                                                                                                                                                                                                                                                                                                                                                                                                                 AAL49057 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JAN-2002; 2002WO-EP000868
                                                                                                                                                                                                     2.9%;
                                                                                                                                                                                                                                                                             265 TGCACCTGGAGCAG 278
                                                                                                                                                                                                                                                                                                                   recaccircaacae 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                       Local Similarity 92.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-636566/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200261121-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Drosophila sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-OCT-2002
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Best Local S
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                                                                                                                                                                                                                                           Matches
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Sequence 18 BP; 6 A; 3 C; 5 G; 4 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating
                                                                                                                                                                                                                                          Human, PRO; benign tumour, malignant tumour; lymphoid malignancy; leukaemia, neuronal disorder; stromal disorder; blastocoelic disorder; inflammatory disorder; immune disorder; angiogenic disorder; cytostatic; neuroprotective; PCR; primer; ss.
                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thirty five nucleic acids encoding PRO polypeptides, useful for benign or malignant tumors, leukemias and lymphoid malignancies. inflammatory, angiogenic and immunologic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hillan KJ;
Stone DM;
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Length 18;
                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gurney
 Score 12.4; DB 1;
Pred. No. 4.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                    Forward PCR primer 4 for human PRO274 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Godowski PJ,
I RM, Roy MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; Page 119; 302pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Goddard A, Godowe
Pan J, Pitti RM,
Wood WI;
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99US-0140650P.
99US-0140653P.
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99US-0145698P.
99US-0146222P.
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99WO-US028301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99WO-US028634
  2.9%;
                                                   230 CAAATCGGGAGGCT 243
                                                                                                                                            ABK40318 standard; DNA; 18
                                                                            CAAAACGGGAGGCT 18
                                                                                                                                                                                             (first entry)
Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GETH ) GENENTECH INC.
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Marsters SA, P
Watanabe CK, W
                                                                                                                                                                                                                                                                                                                                     WO200153486-A1.
                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                       11-FEB-2000;
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02-JUN-1999;
22-JUN-1999;
                                                                                                                                                                                             15-JJL-2002
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                                                                                                                                                                                                                                                                                                                                                               26-JUL-2001
                                                                                                                                                                    ABK40318;
                                                                             'n
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                                                                                                                   RESULT 680
                                                                                                                                ABK40318/c
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18 AAGCTGCTGAAGGA 5

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The present invention describes a method for determining if an individual has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing of a set of target cells, comprising determining if a sample of the subject has an expression product of at least one marker gene. Also described is a compound capable of siglec in a cell. Peripheral blood monouclear cell (PBMC) expressed or Siglec in a cell. Peripheral blood monouclear cell (PBMC) expressed the present invention can be used in a diagnostic method, particularly as an indicator of angiogenesis or to determine presence of a tumour cell. The method of the invention is suitable to determine within a few days if a certain treatment against Kaposi's Sarcoma is successful. Abg81851 to Abg82006 represent nucleotide sequence used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining presence of a tumor cell or angiogenesis, and the effectiveness of treatment, by detecting the presence of marker genes
                                                                                                                                  Gaps
sequences are also useful in gene therapy. The present sequence represents a PCR primer used in the methods of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          effectiveness of treatment, by detecting the presence of mar)
useful to detect and monitor treatment of Karposi's Sarcoma.
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                            Length 18;
                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; Kaposi's sarcoma; tumour; angiogenesis; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 2 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                        Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                          2.9%; Score 12.4; DB 1; 92.9%; Pred. No. 4.3e+02; iive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                   Kaposi's Sarcoma TAG PCR primer SEQ ID NO:142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Page 24; 38pp; English.
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                                                                                                                                                                                                                                                                                                        ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JAN-2001; 2001EP-00200228.
28-SEP-2001; 2001EP-00203703.
28-SEP-2001; 2001US-0325722P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-JAN-2002; 2002EP-00075264.
                                                                                                                                                                      215 GAACTCGGTGGCGG 228
                                                                                                                                                                                                                                                                                                      ABQ81992 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                18 GAACTCCGTGGCGG 5
                                                                                        2.9%
Dest Local Similarity 92.9%
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jan Der Kuyl AC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP1225233-A2.
                                                                                                                                                                                                                                                                                                                                                                                 19-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-JUL-2002.
                                                                                                                                                                                                                                                                                                                                          AB081992;
                                                                                                                                                                                                                                                                RESULT 68
ABQ81992/
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first active agent comprising an oligonucleotide antisense to the first active agent comprising an oligonucleotide antisense to the first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 mucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antistathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antifinialmmatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconscription, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obsained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                    Human, antisense, lung dysfunction, nasal airway dysfunction, antinflammatory steroid, ubiquinone, antinflammatory; antiallergic, antiasthmatic; hypotensive; immunosuppressive, cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Katz E, Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 2 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure, SEQ ID NO 12577; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                Human IL4-R oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nyce JW, Li Y, Sandrasagra A, Ka
Miller S, Tang L, Shahabuddin S;
                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EPIG-) EPIGENESIS PHARM INC
                                      ABZ97335 standard; DNA; 18
                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-229219/22.
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                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                         17-0CT-2003
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                                                                                 ABZ97335;
RESULT
                      ABZ973
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Gaps

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2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; tive 0; Mismatches 1; Indels

282 GGCACCAAGCTGGT 295

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Gaps

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288 AAGCTGGTGAAGGA 301

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13; Conservative

Query Match Best Local Similarity Matches 13; Conserv

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98US-0082804P.
98US-0083736P.
98US-0083332P.
98US-0083332P.
98US-0083332P.
98US-0083496P.
98US-0083496P.
98US-0083496P.
98US-0083446P.
98US-0083584P.
98US-0083584P.
98US-0083584P.
98US-0083584P.
98US-0083584P.
98US-008454P.
98US-008454P.
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98US-008454P.
98US-008454P.
98US-008454P.
98US-008436P.
98US-008454P.
98US-008454P.
98US-008569P.
98US-0091359P.
98US-00168978.
98US-00168978.
98US-00168978.
98US-00168978.
98US-00168978.
98US-00168978.
98US-00168978.
98US-00168978.
98US-00187368.
98US-00187368.
98US-00187368.
99US-002187378.
                                                                                                                                                                                                                                                     Human; secreted and transmembrane protein; PRO; virucide; gene therapy; cell death; growth induction cascade; blood coagulation cascade; viral infection; PCR; primer; 88.
                                                                                                                                                                                                                 Novel human secreted and transmembrane protein related primer #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           970S-0064249P.
970S-0064349P.
970S-0065341P.
980S-0077641P.
980S-0077641P.
980S-0077641P.
980S-0077641P.
980S-0077641P.
980S-0077804P.
980S-0078938P.
980S-0078938P.
980S-0079568P.
980S-007958P.
980S-007958P.
980S-007958P.
980S-007958P.
980S-007958P.
980S-0079338P.
980S-0080137P.
                                                    RESULT 683
ACC42435 Standard; DNA; 18 BP.
XX
XX
XX
C ACC42435;
XX
D1 09-SEP-2003 (first entry)
XX
W Human; secreted and transmemb
XX
XW Human; secreted and transmemb
XX
XW Human; secreted and transmemb
XX
X Home sapiens.
XX
X US2003050239-Al.
XX
X US200303039-Al.
XX
X US200303030303030
                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-OCT-2001; 2001US-00978191
    1 GGCACCAGGCTGGT 14
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Human; secreted and transmembrane protein; PRO; antiinflammatory; antiarteriosclerotic; cardiant; anti-infertility; anti-HIV; cytostatic; antidiabetic; gene therapy; inflammatory disease; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; premature aging; AIDS; cancer; diabetic complication; chromosome mapping; gene mapping; pharmaceutical; diagnostic; biosensor; bioreactor; tissue typing; PCR; primer; ss.
                                                                             Novel human secreted and transmembrane protein related primer #6
                                                                                                                                                                                                                                                                                                                                                                                                       970S-0064249P
970S-00642449P
970S-0066364P
980S-0077452P
980S-0077641P
980S-0077641P
980S-0077641P
980S-0077641P
980S-0077641P
980S-0077641P
980S-007801P
980S-007801P
980S-0079664P
980S-0079786P
980S-0079786P
980S-0079786P
980S-0079786P
980S-0080107P
980S-0080198P
980S-008198P
980S-008198P
980S-008198P
980S-008199P
980S-0082797P
                                                                                                                                                                                                                                                                                                                                                                        24-OCT-2001; 2001US-00999832
                                          16-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                               US2002192706-A1.
                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                   19-DEC-2002
       ACA63470;
       Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Baton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 12.4; DB 1; Length 18; Best Local Similarity 92.9%; Pred. No. 4.3e+02; Matches 13; Conservative 0; Mismatches 1; Indels
14-MAY-1999; 99WO-USO10733.

16-JUN-1999; 99WS-01391252.

16-JUN-1999; 99US-0134268P.

23-JUL-1999; 99US-0134268P.

26-JUL-1999; 99US-0144268P.

28-JUL-1999; 99US-0146222P.

28-JUL-1999; 99US-0146222P.

28-JUL-1999; 99US-0146222P.

25-AUG-1999; 99US-01380142.

25-AUG-1999; 99US-01380142.

25-AUG-1999; 99US-01380142.

25-AUG-1999; 99US-01380142.

25-AUG-1999; 99US-01380142.

30-DEC-1999; 99US-01380142.

30-DEC-1999; 99WO-USO1265.

16-DEC-1999; 99WO-USO1265.

16-DEC-1999; 99WO-USO1365.

11-FEB-2000; 2000WO-USO1365.

11-FEB-2000; 2000WO-USO1365.

11-MAX-2000; 2000WO-USO1365.

11-MAX-2000; 2000WO-USO1365.

22-MAX-2000; 2000WO-USO1365.

23-MAX-2000; 2000WO-USO1365.

24-AUG-2000; 2000WO-USO1365.

25-MAX-2001; 2001WO-USO1365.

25-MAX-2001; 2001WO-USO1365.

10-MAX-2001; 2001WO-USO1365.

11-TPUN-2011; 2001WO-USO1365.

20-UNN-2001; 20
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ACA63470/c
ID ACA63470 standard; DNA; 18
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18
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The invention describes an isolated nucleic acid (I) comprising, or which is at least 80 % sequence identity to, or the full-length coding sequence off, any of 118 300-2100 nucleotide sequences, which encodes its corresponding PRO polypeptide selected from 118 100-700 amino acid sequences, all given in the specification. The nucleic acids and polypeptides are useful for treating inflammatory diseases, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, cancer, or diabetic complications. The nucleic acids are useful as hybridisation probes, in chromosome and gene mapping, and in generating antiseanse RNA or DNA. The polypeptides are useful as pharmaceuticals, diagnostics, blosensors or bioreactors. Both are useful in tissue typing. This sequence represents a novel human secreted and transmembrane PRO polypeptide associated primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New secreted and transmembrane nucleic acids and polypeptides, designated as PRO, useful for treating inflammation, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi UC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napler MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; Page 125; 453pp; English.
20-NOV-1999; 98WO-US024855.
05-JAN-1999; 99WO-US00106.
06-JAN-1999; 99WO-US00106.
10-MAR-1999; 99WO-US001073.
02-JUN-1999; 99WO-US011073.
02-DEC-1999; 99WO-US01252.
30-NOV-1999; 99WO-US01252.
30-DEC-1999; 99WO-US012651.
02-DEC-1999; 99WO-US012651.
03-DEC-1999; 99WO-US012651.
04-JAN-2000; 2000WO-US001219.
05-JAN-2000; 2000WO-US001219.
05-JAN-2000; 2000WO-US001219.
06-JAN-2000; 2000WO-US001219.
06-JAN-2000; 2000WO-US001219.
06-JAN-2000; 2000WO-US001219.
11-PEB-2000; 2000WO-US001319.
11-PEB-2000; 2000WO-US013661.
11-MAR-2000; 2000WO-US001319.
11-MAR-2000; 2000WO-US013765.
11-MAR-2000; 2000WO-US013765.
11-MAR-2000; 2000WO-US013765.
11-MAR-2000; 2000WO-US013705.
17-MAR-2000; 2000WO-US013705.
17-MAR-2000; 2000WO-US013705.
17-MAR-2000; 2000WO-US014914.
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2001WO-US017092.
2001WO-US017800.
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2000WO-US023328
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Score 12.4; DB 1; Length 18; Pred. No. 4.3e+02;

2.9%; 92.9%;

Query Match Best Local Similarity

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                                                                                                                                                                                 Human; ds; thrombolytic agent; interferon; interleukin; cytokine; erythropoietin; colony stimulating factor; cancer; colorectal carcinoma; appprosis related condition; AIDS; amyotrophic lateral sclerosis; inflammatory disease; asthma; atherosclerosis; neurodegenerative disease; gastrointestinal disorder; Alzheimer's disease; Parkinson's disease; hypertension; myocardial ischaemia; Kidney disease; carcinogenesis; plomerulonephritis; lung disease; pulmonary hypertension; preeclampsis; bronchial asthma; gastric ulcer; renal failure; cardiovascular disease; inflammatory bowel disease; reproductive disorder; premature labour.
 Gaps
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                                                                                                                                                                 Human PRO polypeptide associated oligonucleotide SEQ ID NO 14.
 Indels
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 Mismatches
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9703 - 0065311P

9803 - 0077450P

9803 - 0077451P

9803 - 0077641P

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9803 - 0077641P

9805 - 0077650P

9805 - 0078936P

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9805 - 0078656P

9805 - 0079656P
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98US-0079786P.
98US-0079920P.
98US-00105413.
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98US-00184216.
98US-00187368.
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98US-00202054.
98US-00218517.
                                                                                                ACA71634 standard; DNA; 18 BP
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 13; Conservative
                       215 GAACTCGGTGGCGG
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20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
25-MAR-1998;
26-MAR-1998;
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26-JUN-1998;
07-OCT-1998;
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27-MAR-19
27-MAR-19
  Matches
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ACA71634/
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Disclosure, SEQ ID NO 14; 55pp; English.
12-APR-1999; 99US-00284291.
14-MAY-1999; 99US-00311832.
14-MAY-1999; 99WO-US011832.
02-UN-1999; 99WO-US012552.
25-AUG-1999; 99US-00380137.
25-AUG-1999; 99US-00380137.
25-AUG-1999; 99WO-US028513.
02-DEC-1999; 99WO-US028565.
16-DEC-1999; 99WO-US028565.
16-DEC-1999; 99WO-US031243.
30-DEC-1999; 99WO-US031243.
30-DEC-1999; 99WO-US031274.
05-JAN-2000; 2000WO-US000376.
06-JAN-2000; 2000WO-US000376.
11-PEB-2000; 2000WO-US00376.
11-PEB-2000; 2000WO-US00376.
11-PEB-2000; 2000WO-US00376.
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21-MAR-2000; 2000WO-US007332.
30-MAR-2000; 2000WO-US008439.
17-MAY-2000; 2000WO-US013705.
22-MAY-2000; 2000WO-US014042.
30-MAY-2000; 2000WO-US014941.
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2000WO-US020710.
2000WO-US023328.
2000US-00709238.
2000US-00723749.
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2001US-00816920.
2001WO-US009552.
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2001US-00874503.
2001US-00882636.
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22-MAR-2001; 2
22-MAR-2001; 2
10-MAY-2001; 2
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14-JUN-2001;
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Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;

New isolated PRO polypeptides e.g. PRO213, PRO274 and PRO300, for use as pharmaceuticals, diagnostics, biosensors and bioreactors, for identifying modulators of receptor-ligand interactions.

The invention relates to an isolated secreted and transmembrane polypeptide, designated as PRO polypeptide. The PRO polypeptide is useful in PRO polypeptide detection methods. The PRO polypeptide is useful for linking a bioactive molecule to a cell. The PRO polypeptide or an antibody against it is useful for modulating a biological activity of a cell. The PRO polypeptide is useful in industrial applications including

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pharmaceuticals, diagnostics, biosensors and bioreactors. The PRO polypeptide is also useful as a thrombolytic agent, interferon, colory stimulating factor and other cytoropicatin, colony stimulating factor and other cytokines. The PRO polypeptide is useful for treating disease such as cancer e.g. colorectal carcinoma; apoptosis related conditions e.g. AIDS, amyotrophic lateral scleroals; inflammatory disease e.g. Alzheimer's disease, catherosclerosis; neurodegenerative disease e.g. Alzheimer's disease, Parkinson's disease; cardiovascular disease e.g. hypertension and glomerulonephritis; lung disease e.g. pulmonary hypertension and glomerulonephritis; lung disease e.g. pulmonary hypertension, bronchial asthma; gastrointestinal disorders e.g. pulmonary hypertension, bronchial companies is carcinogenesis. The present sequence represents a PRO polypeptide associated oligonucleotide of the invention. Note: The sequence data for this patent did not form part of the printed sequence at sequence. The sequence cate of the invention. Note: The sequence data for this patent did not form part of the printed sequence. The sequence cate of the process of the invention was obtained in electronic format directly from USPTO cat sequence.
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970S-0062250P.
970S-0065311P.
980S-0077632P.
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les 13; Conservative
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Matches
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08-NOV-2000; 2000US-0070228.
27-NOV-2000; 2000US-0072749.
01-DEC-2000; 2000WS-00727759.
20-DEC-2000; 2000WS-00747559.
20-DEC-2000; 2000WS-00747559.
22-MAR-2001; 2001WS-0806520.
22-MAR-2001; 2001US-0081644.
22-MAR-2001; 2001US-00816820.
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2001US-00886342.
2001WO-US019692.
2001WO-US021066.
2001WO-US021735.
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2000WO-US005319
2000WO-US008439
2000WO-US013405
2000WO-US014642
2000WO-US014941
2000WO-US01564
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2001WO-US017800.
2001US-00874503.
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99US-0025465-
99WS-00265686-
99WS-00265686-
99WS-0031831-
99WS-0031831-
99WS-00380138-
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98US-00184216.
98US-00187368.
98WO-USO24855.
98US-00202054.
98US-0079664P-98US-0079664P-98US-0079689P-98US-007928P-98US-0079920P-98US-0079923P-98US-00168978-99US-00168978-
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10-MAY-2001; 2001US-00854280
25-MAY-2001; 2001WO-US017092
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2000WO-US000376.
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27-MAR-1998;
27-MAR-1998;
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07-OCT-1998;
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02-NOV-1998;
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03-MAR-1999;
03-MAR-1999;
05-MAR-1999;
05-MAR-1999;
05-MAR-1999;
06-MAR-1999;
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14-MAY-1999;
14-MAY-1999;
02-JUN-1999;
25-AUG-1999;
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The present invention relates to the isolation of novel human PRO CC polypeptides, and the polymuclectide sequences encoding them. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for inking biological activities of cells expressing PRO polypeptides, for modulating biological activities of cells expressing PRO polypeptides, and for for identifying agonists or antagonists. The bioactive molecule maybe a coxin, radiolabel or antibody, and causes apoptosis or death of the cell. The PRO polypeptides are useful for treating immune disorders, diabetes or hyper or hypo-insulinaemia, cardiac insufficiency, nervous system cisorders, and wound healing. The polymuclectide sequences encoding PRO polypeptides are useful as hybridisation probes, in chromosome and gene compaping, in the generation of antisense RNA and DNA, in the preparation of propagation of antisense RNA and DNA, in the preparation of an immals, for the generation of antisense RNA and DNA, in the preparation of an immals, for the generation sequence represents a PCR primer used in a min in gene therapy. The present invention. Note: The sequence data for this can esqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, 88; PCR; secreted protein; transmembrane protein; PRO; primer; malignancy; cancer; ovarian cancer; colorectal cancer; sarcoma; leukaemia; lymphoma; inflammatory disease; necrosis; atherosclerosis; infertility; premature aging; psortiasis; inflammatory disease; renal disease; arthritis; immune-mediated alopecia; stroke; encephalitis; hepatitis; multiple sclerosis; gene therapy.
Ashkenazi A, Baker KP, Botstein D, Desnoyers L, Eaton D;
Perrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi UC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                  Novel secreted and transmembrane polypeptides and polynucleotides encoding them useful for treating cancer, kidney diseases, bone, cartilage disorders and immune deficiencies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; arive 0; Mismatches 1; Indels
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Best Local Similarity 92.9
Matches 13, Conservative
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03-NOV-1997;
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97105 - 0066341P.
97105 - 0066344P.
98105 - 0077643P.
98105 - 0078044P.
98105 - 0078044P.
98105 - 0078044P.
98105 - 0078044P.
98105 - 007924P.
98105 - 0079728P.
99105 - 0079728P.

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2000WO-US034956
2001WO-US006520
  13-NOV-1997;
11-NAR-1998;
11-NAR-1998;
11-MAR-1998;
12-MAR-1998;
12-MAR-1998;
20-MAR-1998;
21-MAR-1998;
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21-MAR-1999;
21-MAR-1999;
22-DEC-1999;
22-DEC-1999;
22-MAR-1999;
30-MAR-1999;
30-MAR-1998;
30-MA
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22 MAR. 2001; 2010US-00816920.
22 MAR. 2001; 2010US-00816920.
10 MAY. 2001; 2001US-00854280.
10 MAY. 2001; 2001US-00854280.
25 MAY. 2001; 2001US-00854280.
01 JUN 2001; 2001US-00872035.
01 JUN 2001; 2001US-00882636.
19 JUN 2001; 201US-00882636.
19 JUN 2001; 201US-00882636.
19 JUN 2001; 201US-008862636.
19 JUN 2001; 201US-00886636.
20 JUN 2001; 201US-00886636.
30 JUL 2001; 2001US-00886636.
2001US-00816744.
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### (GETH ) GENENTECH INC.

Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Goddowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kwo SS, Napter MA, Pan J, Paoni NF, Roy MA, Shelton Stewart TA, Tumas D, Williams PM, Wood WI;

#### WPI; 2003-341189/32.

New genes and secreted and transmembrane polypeptides (e.g. PRO337 or PRO1559), useful for treating or diagnosing e.g. cancers, atherosclerosis, infertility, stroke, encephalitis, hepatitis or multiple sclerosis in mammals.

## Example 4; Page 121; 460pp; English.

The invention relates to a new isolated nucleic acid molecule comprises a gequence with at least 80% identity to: (a) a nucleotide encoding any of 94 multiple of 94 pRO polypeptides whose sequences are fully defined in the specification; or the full length coding sequence of any these 94 mucleotide sequences. Also included are an isolated PRO polypeptide corrupts at least 80% politypeptide are an isolated PRO polypeptide having at least 80% amino acid sequence dream isolated PRO polypeptide having corring at least 80% amino acid sequence identity to: (a) an amino acid sequence corrupts at least 80% amino acid sequence identity to: (a) an amino acid sequence corrupts at least 80% amino acid sequence identity to: (a) an amino acid sequence corrupts associated signal specification; (b) the PRO polypeptide, lacking its associated signal peptide, avector comprising the nucleic peptide; or (c) an extracellular domain of the PRO polypeptide, with or lacking its associated signal peptide), a vector comprising the nucleic peptide; or (a) a chimaeric molecule comprising the vector (and polypeptide fused corrupts and an anin-PRO acid sequence and an anti-PRO acid sequence or polypeptides are useful as pharmaceuticals, inclammatory disease, diagnostics, biosensors or bioreactors. These are particularly useful for detecting or treating e.g. malignancies or cancers (e.g. ovarian cancer, colorectal cancer, sarcoma, leuksemia or lymphoma), inflammatory disease, infertility, premature adjing, psoriasis, infertility, premature adjing, psoriasis, infertility, premature, adjing, psoriasis, infertility, premature, adjing, psoriasis, infertility, are propappides are useful as molecular weight markers, or for chromosome corrupts, and these diseases. The PRO polypeptides are useful as molecular weight markers, or for chromosome calso useful as molecular weight markers, or for chromosome calso useful as molecular weight markers, or for chromosome calso the present sequence is a per per infer energy, particularly for replacing a propoly

# Sequence 18 BP; 2 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

Gaps ; 0 Ouery Match
2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels

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rng.res

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Human; secreted and transmembrane protein; PRO; tissue typing; chromosome identification; vaccine; cancer; retinal disorder; sports-related joint disorder; osteoarchitis; rheumatoid arthritis; wound healing; obesity; diabetes; hearing loss, second cardiac insufficiency disorder; kidney disorder; nervous system disorder; haemoglobin associated disorder; PCR; primer; ss.
                                                                                           Secreted and transmembrane PRO protein associated primer #8
                                                                                                                                                                                                                            970S-0062250P-
970S-0064249P-
970S-0066341P-
980S-0077461P-
980S-0077642P-
980S-0077642P-
980S-0077642P-
980S-0077642P-
980S-0077642P-
980S-0077642P-
980S-00780294P-
980S-00780294P-
980S-0079668P-
980S-007968P-
980S-007968P-
980S-0079723P-
980S-0079723P-
980S-0080194P-
980S-0081049P-
980S-0081049P-
980S-0081839P-
980S-0081049P-
                                             ADA24553 standard; DNA; 18 BP
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                                                                          20-NOV-2003 (first entry)
18 GAACTCCGTGGCGG
                                                                                                                                                                                US2003050241-A1.
                                                                                                                                                                 Homo sapiens.
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11. MAR. 1998

11. MAR. 1998

12. MAR. 1998

20. MAR. 1998

20. MAR. 1998

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20. MAR. 1998

25. MAR. 1998

27. MAR. 1998

21. MAR. 1998

31. MAR. 1998
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                                                             ADA24553;
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PR 22-APR-1998; 98US-0082704P. PR 22-APR-1998; 98US-0082795P. PR 22-APR-1998; 98US-0082326P. PR 29-APR-1998; 98US-0083322P. PR 29-APR-1998; 98US-0083322P. PR 29-APR-1998; 98US-0083495P. PR 22-APR-1998; 98US-0083339P. PR 23-APR-1999; 98US-0133022P. PR 23-APR-1999; 98US-0133022P. PR 23-APR-1999; 98US-0133022P. PR 23-APR-1999; 98US-0133022P. PR 23-APR-1999; 99US-0133022P. PR 23-APR-1999; 99US-0133022P.
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2001US-00978403
                                                                                                                              (first entry)
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27-MAR-1998;
30-MAR-1998;
30-MAR-1998;
31-MAR-1998;
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                                                                                                                              08-SEP-2003
                                           ACD29616;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes an isolated secreted and transmembrane (PRO) polypeptide (I). PRO317 polypeptide is useful for detecting PRO4993 polypeptide in a sample, and vice versa. PRO725, PRO700 and PRO739 are useful for detecting PRO725, PRO700 and PRO739 in a sample, and PRO1559 is useful for detecting PRO725, PRO700 and PRO739 in a sample, PRO4993 is polypeptide, and PRO337 is useful for linking a bioactive molecule to a cell expressing a PRO337 polypeptide, PRO705 is useful for linking a bioactive molecule to a cell expressing a PRO337 polypeptide, PRO705, PRO700 and PRO739 polypeptide, PRO705, PRO700 and PRO739 polypeptide, and PRO735, PRO700 and PRO739 polypeptide, and PRO735, PRO700 and PRO739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KP, Botstein D, Desnoyers L, Eaton DL;
E, Fong S, Gao W, Gerber H, Gerritsen ME;
PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Walpier MA, Pan J, Paoni NP, Roy MA, Shelto Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 132; 461pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-AUG-2000; 2000WO-US023328.
01-DEC-2000; 2000WO-US032678.
20-DEC-2000; 2000WO-US034556.
22-MAR-2001; 2001WO-US006520.
22-MAR-2001; 2001WO-US009552.
25-MAY-2001; 2001WO-US009552.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUN-2001; 2001WO-US017800
20-JUN-2001; 2001WO-US019692
29-JUN-2001; 2001WO-US021066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-JUL-2001; 2001MO-US021735
30-JUL-2001; 2001US-00918585
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Filvaroff E, I
Godowski PJ, C
Kuo SS, Napie,
Tumas D, Will,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAACTCCGTGGCGG 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-521814/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
30-NOV-1999;
02-DEC-1999;
16-DEC-1999;
30-DEC-1999;
30-DEC-1999;
30-DEC-1999;
05-JAN-2000;
06-JAN-2000;
11-FEB-2000;
11-FEB-2000;
21-MAR-2000;
21-MAR-2000;
21-MAR-2000;
21-MAR-2000;
22-MAY-2000;
22-MAY-2000;
22-MAY-2000;
30-MAY-2000;
23-MAY-2000;
24-MAY-2000;
24-MAY-2000;
25-MAY-2000;
26-MAY-2000;
26-MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ashkenazi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Matches
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RESULT 689 ACD29616/c ID ACD29616 standard; DNA; 18 BP.

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9805-0083499P9805-0083560P9805-0083558P9805-0083558P9805-0083558P9805-0083742P9805-0084414P9805-0084411P9805-0084631P9805-0084631P-

28-APR-1598;
29-APR-1598;
29-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
29-APR-1998;
30-APR-1998;
31-APR-1998;

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The invention describes an isolated, secreted and transmembrane polypeptide, termed PRO polypeptide (I). (I) is useful for detecting PRO4993, PRO329, PRO725, PRO700 or PRO739 polypeptide, and for linking a bioactive molecule to a cell expressing the above polypeptides. The bioactive molecule is a toxin, radiolabel or an antibody and causes cell death. (I) is useful as therapeutic agent, in medical and industrial applications e.g. for treating neuropathy, especially peripheral neuropathy, dispecially peripheral neuropathy, discassociated neuropathy, charcot-Marie-Tooth disease, Refusum's disease, Abetalipoproteinaemia, Tangier disease, Krabbe's disease, Metachromatic leukodystrophy, Pabry's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME;
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF, Roy MA, Shelton DL;
Stewart TA, Tumas D, Williams PM, Wood WL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel secreted and transmembrane polypeptide for modulating biological activity of cell expressing the polypeptide, identifying agonists or antagonists of polypeptide, and as molecular weight markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       #4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane polypeptide PRO274 primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; Page 125; 459pp; English.
              06-JAN-2000; 2000WO-US000277.

06-JAN-2000; 2000WO-US000376.

11-FEB-2000; 2000WO-US003565.

18-FEB-2000; 2000WO-US0036131.

24-FEB-2000; 2000WO-US005811.

10-MAR-2000; 2000WO-US006811.

11-MAY-2000; 2000WO-US006813.

17-MAY-2000; 2000WO-US014941.

17-MAY-2000; 2000WO-US014941.

17-MAY-2000; 2000WO-US014941.

17-MAY-2000; 2000WO-US014941.

17-MAY-2000; 2000WO-US014961.

18-AUG-2000; 2000WO-US015567.

24-AUG-2000; 2000WO-US015650.

24-AUG-2000; 2000WO-US015650.

25-MAR-2001; 2001WO-US003552.

25-MAR-2001; 2001WO-US003652.

25-MAR-2001; 2001WO-US001560.

26-JUN-2001; 2011WO-US017800.

29-JUN-2001; 2011WO-US017800.

29-JUN-2001; 2011WO-US017800.

29-JUN-2001; 2011WO-US017860.
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ID ADA12214 standard; DNA; 18 BP
XX
AC ADA12214;
XX
DT 06-NOV-2003 (first entry)
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GETH ) GENENTECH INC.
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98US-0084640P 98US-0084643P 98US-0085338P 98US-0085338P 98US-0085339P 98US-0085339P 98US-008589P 98US-008648P 98US-01088P 99US-011361P 99US-011362P 99US-011365P 99US-011365P 99US-011368P 99US-011368P 99US-011368P

12-MAR-1999; 29-MAR-1999; 21-APR-1999; 26-APR-1999;

99WO-US031243 99WO-US031274

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30 - APR - 1998 |
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14 - MAR - 1
  primer; ss; inflammatory disease; organ failure; atherosclerosis; cardiac injury; infertility; birth defect; premature aging; AIDS; cancer; diabetic complication; tissue typing; human; PCR.
                                                                                                                                     9504S-0018049P-
9710S-0056226P-
9710S-0056236P-
9710S-00563311P-
9810S-00774549P-
9810S-00774541P-
9810S-00777431P-
9810S-0077731P-
9810S-0077731P-
9810S-0078931P-
9810S-0080333P-
9810S-0081197P-
9810S-0081333P-
9810S-0081333P-
9810S-00813332P-
9810S-0083332P-
                                                                                                                2001US-00978824
                                                                    US2003055216-A1
                                                                                                                                     21-MAY-1996;
17-007-1997;
13-NOV-1997;
13-NOV-1997;
11-MAR-1998;
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12-MAR-1998;
13-MAR-1998;
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31-MAR-1998;
32-MAR-1998;
31-MAR-1998;
32-MAR-1998;
31-MAR-1998;
32-APR-1998;
32-APR-1998;
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32-APR-1998;
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29-APR-1998;
29-APR-1998;
29-APR-1998;
                                               Homo sapiens
                                                                                          20-MAR-2003
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98US-0083742P.
98US-0084436EP.
98US-0084431P.
98US-0084431P.
98US-0084530P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0084637P.
98US-0086373P.
98US-0085637P.
98US-0085637P.
98US-0085637P.
98US-008563P.
98US-00863P.
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98US-008563P.
98US-00863P.
98US-013253P.
99US-013353P.
99US-013353P.
99US-0133537P.

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diabetic peripheral neuropathy; autonomic neuropathy; reduced motility of the gastrointestinal tract; atony of the uniary bladder; post polio syndrome; Krabbe's disease; Charcot Marie-Tooh disease; Fabry's disease; Tangier disease; Refsum's disease; Papry's disease; Tangier disease;
cardiac- insufficiency disorder; peripheral neuropathy;
                                                                                                                                                                                                               97US-0064249P.
97US-0064449P.
97US-0065844P.
97US-0065844P.
98US-0077641P.
98US-0077641P.
98US-0077641P.
98US-0077641P.
98US-0070894P.
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98US-0070898P.
98US-0070898P.
98US-0070988P.
98US-0080107P.
98US-0080107P.
98US-0080134P.
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98US-0080134P.
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                                                                                                                              US2003049633-A1.
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29-APR-1998;
29-APR-1998;
                                                                                                     Homo sapiens.
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                                                                                                                                                            13-MAR-2003
 Human; secreted and transmembrane protein; PRO; viral infection; tumour growth, retinal disorder; injury; sight loss; retinitis pigmentosum; age-related macular degeneration; sport-related joint problem; articular cartilage defect; osteoarthritis; rheumatoid arthritis; wound healing; obesity; diabetes; insulinaemia; kidney disorder; mesangial cell function; Berger disease; nephropathy; celiac disease; dermatitis; Crohn disease; neuropathy;
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Fong S, Gao W, Gerber H, Gerritsen
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
                                                                                                                18-FEB-2000; 2000MC-0S004341.
24-FEB-2000; 2000MC-0S004341.
10-MAR-2000; 2000MC-0S005819.
21-MAR-2000; 2000MC-0S005819.
21-MAR-2000; 2000MC-0S005819.
30-MAR-2000; 2000MC-0S014042.
30-MAY-2000; 2000MC-0S014042.
30-MAY-2000; 2000MC-0S014042.
30-MAY-2000; 2000MC-0S014042.
30-MAR-2000; 2000MC-0S014041.
28-JUL-2000; 2000MC-0S0126441.
28-JUL-2000; 2000MC-0S0129141.
28-JUL-2000; 2000MC-0S0129141.
28-JUL-2000; 2000MC-0S0129141.
28-JUL-2000; 2000MC-0S0129141.
28-JUL-2000; 2000MC-0S0129141.
28-JUL-2000; 2000MC-0S01491.
28-JUL-2000; 2000MC-0S0129141.
22-MAR-2001; 2001WC-0S016920.
22-MAR-2001; 2001WC-0S016920.
32-MAR-2001; 2001WC-0S016920.
32-MAR-2001; 2001WC-0S016920.
32-MAR-2001; 2001WC-0S016920.
31-MAY-2001; 2001WC-0S016920.
31-MAY-2001; 2001WC-0S017092.
 02-DEC-1999; 99WO-USO28565.
16-DEC-1999; 99WO-USO31095.
30-DEC-1999; 99WO-USO31243.
30-DEC-1999; 99WO-USO31243.
30-DEC-1999; 99WO-USO31274.
65-JAN-2000; 2000WO-USO00277.
66-JAN-2000; 2000WO-USO00376.
11-FEB-2000; 2000WO-USO0376.
11-FEB-2000; 2000WO-USO0376.
11-FEB-2000; 2000WO-USO0376.
11-FEB-2000; 2000WO-USO0376.
11-MAR-2000; 2000WO-USO05319.
11-MAR-2000; 2000WO-USO19705.
11-MAR-2000; 2000WO-USO19705.
11-MAR-2000; 2000WO-USO19706.
11-MAR-2000; 2000WO-USO19706.
11-MAR-2000; 2000WO-USO19706.
11-MAR-2000; 2000WO-USO19706.
12-MAY-2000; 2000WO-USO19706.
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2001US-00882636.
2001US-00886342.
2001WO-US019692.
2001WO-US021066.
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Ferrara N, Filvaroff E,
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ACD29031/
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98US-0083554P.
98US-0083558P.
98US-0083558P.
98US-008456P.
98US-0084541P.
98US-0084637P.
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98US-0084637P.
98US-0085838P.
98US-0085838P.
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98US-0085838P.
98US-0085838P.
98US-0085838P.
98US-008583P.
98US-013322P.
99US-013323P.
99US-01313287.
99US-01313287.
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99US-01313287.
99US-01313287.
99US-01313287.
29-APR-1998;
29-APR-1998;
30-APR-1998;
30-APR-1998;
06-MAY-1998;
06-MAY-1998;
07-MAY-1998;
07-MAY-1998;
07-MAY-1998;
07-MAY-1998;
07-MAY-1998;
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22-MAY-1998;
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22-MAY-1998;
23-MAY-1998;
26-UUN-1998;
26-UUN-1998;
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12-MAR-1999;
29-MAR-1999;
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25-AUG-1999
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Gaps
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29-OCT-1999; 99WS-0162506P.

20-DEC-1999; 99WO-US028513.

02-DEC-1999; 99WO-US028551.

02-DEC-1999; 99WO-US028551.

02-DEC-1999; 99WO-US031035.

30-DEC-1999; 99WO-US0310243.

30-DEC-1999; 99WO-US031274.

30-MAX-2000; 2000WO-US03131.

30-MAX-2000; 2000WO-US03131.

30-MAX-2000; 2000WO-US03131.

30-MAX-2000; 2000WO-US03131.

30-MAX-2000; 2000WO-US03131.

30-MAY-2000; 2000WO-US03131.

30-MAY-2000; 2000WO-US03131.

30-MAY-2000; 2000WO-US03131.

30-DEC-2000; 2000WO-US0313749.

31-DEC-2000; 2000WO-US0313749.

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AU-DEC-2003 (first entry)

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16-OCT-2001; 2001US-00978608

US-0062250P US-0064249P US-0065311P US-0065341P US-0077632P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-0077641P US-007861P US-007861P	US-00796659PUGS-00796639PUGS-00796639PUGS-00796639PUGS-0079728PUGS-0079969PUGS-0079969PUGS-0079969PUGS-0080107PUGS-0080195PUGS-0080195PUGS-0080197PUGS	98US-0082568P. 98US-0082704P. 98US-0082704P. 98US-0082704P. 98US-0082797P. 98US-0083795P. 98US-0083332P. 98US-0083332P. 98US-0083495P. 98US-0083495P. 98US-0083495P. 98US-0083495P. 98US-0083554P. 98US-0083554P. 98US-0084441P. 98US-0084650P. 98US-0084650P. 98US-0084653P. 98US-0084653P.
- CCT - 1997 - NOV - 1997 - NOV - 1997 - NOV - 1998 - MAR - 1998		21-APR-1998; 22-APR-1998; 22-APR-1998; 22-APR-1998; 22-APR-1998; 23-APR-1998; 24-APR-1998; 29-APR-1998; 30-APR-1998; 31-APR-1998; 31-APR-1998; 31-APR-1998;
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- MAX - 1 - MAX	26-JUN-199 226-JUN-199 226-JUN-199 226-JUN-199 21-JUL-199 23-JUL-199 23-JUL-199 20-JUN-1		6-DEC- 0-DEC- 0-DEC- 0-DEC- 0-DEC- 6-JAN- 6-JAN- 1-FEB- 8-FEB- 2-MAR- 0-MAR-
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Gaps
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 21-MAR-2000; 2000WO-US007532.
30-MAY-2000; 2000WO-US018439.
12-MAY-2000; 2000WO-US0184042.
22-MAY-2000; 2000WO-US014042.
30-MAY-2000; 2000WO-US014041.
30-MAY-2000; 2000WO-US015264.
28-UUL-2000; 2000WO-US015264.
28-UUL-2000; 2000WO-US015264.
28-UUL-2000; 2000WO-US015264.
28-UUL-2000; 2000WO-US015264.
20-MCC-2000; 2000WS-00723749.
30-DEC-2000; 2000WS-00723749.
30-MAR-2001; 2001WS-08016674.
32-MAR-2001; 2001WS-00816674.
32-MAR-2001; 2001WS-00816674.
32-MAR-2001; 2001WS-00816678.
32-MAY-2001; 2001WS-0081678.
32-MAY-2001; 2001WS-00814508.
32-MAY-2001; 2001WS-00814508.
32-MAY-2001; 2001WS-00814508.
31-UUN-2001; 2001WS-00816335.
31-UUN-2001; 2001WS-00816335.
31-UUN-2001; 2001WS-00816318.
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97US-0064249P.
97US-0065311P.
97US-0077450P.
98US-0077632P.
98US-0077612P.
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03-NOV-1997;
13-NOV-1997;
21-NOV-1997;
11-MAR-1998;
11-MAR-1998;
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Best Local S
Matches 13
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98US-0086023P. 98US-0086392P. 98US-0086414P.

15-MAY-1998; 18-MAY-1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 22-MAY-1998; 23-MAY-1998; 24-MAY-1998; 26-JUN-1998; 26-JUN-1998; 26-JUN-1998; 20-JUL-1998; 20-NOV-1998; 22-DEC-1998;

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The present invention relates to the isolation of novel human PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are secreted and transmembrane proteins. The PRO polypeptides are useful for detecting other PRO polypeptides, for linking bioactive molecules to cells expressing PRO polypeptides, for modulating biological activities of cells expressing PRO polypeptides, and for toxin, radiolabel or antibody, and cause cell death. The PRO polypeptides are useful for treating neuropathy and neuropathy related diseases such as Charcher-Marie-Tooth disorder, Refoun's diseases, and Krabbe's disease. The polynucleotide sequences encoding PRO polypeptides are useful as The polynucleotide sequences encoding PRO polypeptides are useful as hybridisation probes, in chromosome and gene mapping, in the generation of antisense RNA and DNA, in the preparation of PRO polypeptides, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, 88; PCR, secreted protein, transmembrane protein, PRO; cytostatic, ophthalmological, antiarthritic, osteopathic, antirheumatic, vulnerary, auditory, tumour growth, retinal disorder; sports-related joint problem, articular cartilage defects, osteoarthritis, rheumatoid arthritis; wound healing, hearing loss; primer.
                                                                                       New PRO polypeptides useful for treating peripheral neuropathy, neuropathies associated with systemic disease such as post-polio syndrome or AIDS-associated syndrome.
 Hillan KJ;
Roy MA, Shelton DL;
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2.9%; Score 12.4; DB 1; Length 18;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1; Indels
Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, 1
Kljavin IJ, Kuo SS, Napier MA, Pan J, Paoni NF,
Stewart TA, Tumas D, Williams PM, Wood WI;
                                                                                                                                                              Example 4; Page 125; 425pp; English.
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97US-0064249P.
97US-006531IP.
98US-0077450P.
98US-0077641P.
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13-NOV-1997;
10-MAR-1998;
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Gerritsen ME;
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Desnoyers L, W, Gerber H,

Botstein D, Fong S, Gao

f, Baker KP, Filvaroff E,

Ashkenazi AJ, Ferrara N, F

GETH ) GENENTECH INC.

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24-FEB-2000; 2 02-MAR-2000; 2 10-MAR-2000; 2 21-MAR-2000; 2 30-MAR-2000; 2

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05-JAN-1999) 10-MAR-1999) 10-MAR-1999) 29-MAR-1999) 21-APR-1999) 21-APR-1999) 21-APR-1999) 21-APR-1999) 22-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999) 23-JUN-1999)

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PR 27-WAR.1998; 98US-0079664P.
PR 27-WAR.1998; 98US-0079664P.
PR 27-WAR.1998; 98US-0079786P.
PR 21-WAR.1998; 98US-0080137P.
PR 01-APR-1998; 98US-0080137P.
PR 01-APR-1998; 98US-0080137P.
PR 01-APR-1998; 98US-008123P.
PR 15-APR-1998; 98US-008123P.
PR 22-APR-1998; 98US-008123P.
PR 22-APR-1998; 98US-008132P.
PR 23-APR-1998; 98US-008133P.
PR 23-APR-1998; 98US-008133P.
PR 23-APR-1998; 98US-008133P.
PR 23-APR-1998; 98US-00
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PR 22-MAY-1998; 98US-0086486 P. PR 28-MAY-1998; PR 28-MAY-1999; PR 28-MAY-1999

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22-MAR-2001; 2001US-00B16744.
22-MAR-2001; 2001US-00B16920.
22-MAR-2001; 2001US-00B562.
10-MAY-2001; 2001US-00B5420B.
10-MAY-2001; 2001US-00B5420B.
10-MAY-2001; 2001US-00B5420B.
01-JUN-2001; 2001WS-00B7035.
01-JUN-2001; 2001US-00B7535.
19-JUN-2001; 2001US-00B82635.
20-JUN-2001; 2001WS-0US019692.
20-JUN-2001; 2001WS-US019692.
20-JUN-2001; 2001WS-US019692.
20-JUN-2001; 2001WS-US01966.
09-JUL-2001; 2001WS-US019653.
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98US-0078939P
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98US-0079558P
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18 GAACTCCGTGGCGG 5
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11. SEP-1998; 98US-0100338PF. O7-OCT-1998; 98US-0100338PF. O1-OCT-1998; 98US-0100338PF. O1-OCT-1998; 98US-0100338PF. O1-DCT-1998; 98US-01018748P. O2-NOV-1998; 98US-01018748PF. O2-NOV-1998; 98US-01018748PF. O2-NOV-1998; 98US-01018748PF. O2-NOV-1998; 98US-0101824PF. O2-NAR-1999; 98US-013296PF. O2-NAR-1999; 98US-013296PF. O2-NAR-1999; 99US-013296PF. O2-NAR-1999; 99US-013296PF. O2-NAR-1999; 99US-013296PF. O2-NAR-1999; 99US-0130232PF. O2-NAR-2000; 2000WO-US00231243. O2-NAR-2000; 2000WO-US00380131. O2-NAR-2000; 2000WO-US00380132. O2-NAR-2000; 2000WO
98US-0094651P.
98US-0100038P.
98US-0100038P.
98US-01084216.
98US-01084216.
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98US-01084216.
98US-01084216.
98US-01084285.
98US-010802686.
99US-01080528.
99US-01080528.
99US-01080528.
99US-01080528.
99US-0134281P.
99US-0134281P.
99US-0134287P.
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Human, ss; PCR, secreted protein; transmembrane protein; PRO; cytostatic; ophthalmological; antiarthritic; osteopathic; antirheumatic; vulnerary; auditory; tumour growth, retinal disorder; sports-related joint problem; articular cartilage defects; osteoarthritis; rheumatoid arthritis; wound healing; hearing loss; primer.
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2.9%; Score 12.4; DB 1;
Best Local Similarity 92.9%; Pred. No. 4.3e+02;
Matches 13; Conservative 0; Mismatches 1;
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05-JUN-2001; 2001US-00874503.
14-JUN-2001; 2001US-00882636.
19-JUN-2001; 2001US-0086542.
20-JUN-2001; 2001WO-US021066.
29-JUN-2001; 2001WO-US021066.
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PR 20-NOV-1998; 980G-0103304P; PR 20-NOV-1998; 980G-010324855. PR 22-DEC-1998; 980G-010324855. PR 22-DEC-1998; 980G-010324855. PR 22-DEC-1998; 980G-01032654. PR 22-DEC-1998; 980G-010326285. PR 23-DEC-1999; 980G-010326285. PR 10-WAR-1999; 980G-010326286. PR 10-WAR-1999; 980G-010326287. PR 10-WAR-1999; 980G-010326287. PR 12-WAR-1999; 980G-01030237. PR 12-WAR-1999; 990G-01030237. PR 12-WAR-2000; 2000WO-010009565. PR 12-WAR-2000; 2000WO-010099565. PR 12-WAR-2000; 2000WO-010099565. PR 12-WAR-2000; 2000WO-0100099565. PR 12-WAR-2000; 2000WO-010099565. PR 12-WAR-2001; 2001WO-010099565. PR 12-WAR-2001; 2001WO-010099665. PR 12-WAR-2001; 2001WO-010099695. PR 12-WAR-2001; 2001WO-010099695. PR 12-WAR-2001; 2001WO-

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                                                   2.9%; Score 12.4; DB 1; Length 18; 92.9%; Pred. No. 4.3e+02; tive 0; Mismatches 1; Indels
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970S-0064249P
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980S-0079664P
30-JUL-2001; 2001US-00918585.
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                                                                                              215 GAACTCGGTGGCGG 228
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                                                                                                                   18 GAACTCCGTGGCGG 5
                                                              Best Local Similarity 92.9
Matches 13; Conservative
                     (GETH ) GENENTECH INC.
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                                                      Query Match
                                                                                                                                                    RESULT 697
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Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Baton DL; Ferrara N, Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME; Goddard A, Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ; Kljavin IJ, Kwo SS, Napier MP, Pan J, Paoni NF, Roy MA, Shelton DL; Stewart TA, Tumas D, Williams PM, Wood WI;
05-JAN-1999, 99W0-USD00106.
08-MAR-1999, 99W5-00254465.
08-MAR-1999, 99W5-002656861.
10-MAR-1999, 99W0-USD0190.
12-AR-1999, 99W0-USD0190.
12-AR-1999, 99W0-USD0190.
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14-MAR-1999, 99W0-USD01913.
15-AR-2000, 2000W0-USD019565.
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11-TEB-2000, 2000W0-USD019565.
11-TEB-2000, 2000W0-USD019565.
11-MAR-2000, 2000W0-USD01991.
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13-AR-2000, 2000W0-USD01991.
13-AR-2001, 2000W0-USD01991.
13-AR-2001, 2000W0-USD01991.
13-AR-2001, 2000W0-USD01992.
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10-MAY-2001; 2001US-00854208.

25-MAY-2001; 2001US-00854280.

25-MAY-2001; 2001US-00854208.

10-UNA-2001; 2001US-008762035.

10-UNA-2001; 2001US-00876503.

14-UNA-2001; 2001US-00886542.

20-UNA-2001; 2001US-00886542.

20-UNA-2001; 2001US-00886592.

20-UNA-2001; 2001US-00886592.

20-UNA-2001; 2001US-00886592.

20-UNA-2001; 2001US-00886592.
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Novel secreted and transmembrane polypeptides and polynucleotides accoding them, useful for treating wound healing, tissue growth and muscle generation and regeneration, amyotrophic lateral sclerosis or neuropathy.

WPI; 2003-596568/56.

Example 4; SEQ ID NO 14; 472pp; English.